

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 109354

TO: Terra Gibbs

Location: CM1/12A12/11E12

Art Unit: 1635

Monday, December 01, 2003

Case Serial Number: 09/954556

From: David Schreiber

Location: Biotech-Chem Library

CM1-6A03

Phone: 308-4292

david.schreiber@uspto.gov

Search Notes	
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SEARCH REQUEST FORM

	Requestor's Name:		Serial Number: _	
	Date:	Phone:		Art Unit:
·	Search Topic: Please write a detailed statement of search terms that may have a special meaning. Giplease attach a copy of the sequence. You is	ve examples or relevent citat	ions, authors, l	subject matter to be searched. Define any keywords, etc., if known. For sequences, nost relevent claim(s).
	Date completed: 12/1 Searcher: 9.5 (Nor 200 30) Terminal time: 93 Elapsed time: 17 CPU time: Total time: Number of Searches: Number of Databases:	C P Type of S N A Si	te TIC M-1 640 re-S	Vendors IG STN Dialog APS Geninfo SDC DARC/Questel U Other G in pages

PTO-1590 (9-90)

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From:

Gibbs, Terra

Sent:

Friday, November 21, 2003 12:14 PM

To:

Schreiber, David

Subject:

Sequence search request...

Hi David,

Doug Schultz and Karen LaCourcicie recommended that I send you this search request.

I have a request for a score over length search:

I need a length limited nucleotide sequence search against nucleobases 1317-2720 of SEQ ID NO:3 of USSN 09/954,556, where the returns are rank ordered based on the score over length/ratio as we've discussed. I need the lengths limited to hits between 8 and 50 nucleotides, and I'll take as many hits as you can import into excel (64,000?), and alignments for anything above .75 on the above ratio. Hope this is clear, please call me if it's not.

Terra Cotta Gibbs, Ph.D. Art Unit 1635 CM1, 12A12 703-306-3221

A E - 4.4

OLDON MANAGERA SINT

Schreiber, David

From:

Fredman, Jeffrey

Sent:

Tuesday, November 25, 2003 7:06 AM

To: Cc: Schreiber, David Gibbs, Terra

Subject:

FW: Sequence search request (RUSH)...

PLEASE RUSH

I Approve.

Jeff Fredman

-----Original Message-----

From:

Gibbs, Terra

Sent:

Monday, November 24, 2003 10:40 AM

To:

Fredman, Jeffrey

Subject:

FW: Sequence search request (RUSH)...

This case is an AF. I need an immediate sequence search.

Could you please approve the RUSH?

Also, this sequence search request has to be sent to David Schreiber, instead of STIC-Biotech in general.

Thanks.

-----Original Message-----

From:

Gibbs, Terra

Sent:

Friday, November 21, 2003 12:14 PM

To:

Schreiber, David

Subject:

Sequence search request...

Hi David,

Doug Schultz and Karen LaCourcicie recommended that I send you this search request.

I have a request for a score over length search:

I need a length limited nucleotide sequence search against nucleobases 1317-2720 of SEQ ID NO:3 of USSN 09/954,556, where the returns are rank ordered based on the score over length/ratio as we've discussed. I need the lengths limited to hits between 8 and 50 nucleotides, and I'll take as many hits as you can import into excel (64,000?), and alignments for anything above .75 on the above ratio. Hope this is clear, please call me if it's not.

Terra Cotta Gibbs, Ph.D. Art Unit 1635 CM1, 12A12 703-306-3221 OHEN MATERIAL SILL



STIC SEARCH RESULTS FEEDBACK FORM

Biotech-Chem Library

Questions about the scope or the results of the search? Contact the searcher or contact:

Mary Hale, Information Branch Supervisor 308-4258, CM1-1E01

Voluntary Results Feedback Form
> I am an examiner in Workgroup: Example: 1610
> Relevant prior art found , search results used as follows:
☐ 102 rejection
☐ 103 rejection
☐ Cited as being of interest.
Helped examiner better understand the invention.
Helped examiner better understand the state of the art in their technology.
Types of relevant prior art found:
☐ Foreign Patent(s)
Non-Patent Literature (journal articles, conference proceedings, new product announcements etc.)
> Relevant prior art not found:
Results verified the lack of relevant prior art (heiped determine patentability).
Results were not useful in determining patentability or understanding the invention.
Comments

Disposition and completed forms to STICE to the Circum Library Gilli — Circ. Desk



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RESULT 2 \$4135581 LOCUS BEK=fibroblast growth factor receptor { leukocyte DNA, Genomic, 36 nt, segment } ACCESSION \$41355.1 GI:232802 KEYMORDS SEGMENT 1 of 2 SOURCE Homo sapiens (human) ORGANISM Eukaryota; Metazoa; Chordata; Craniata;	Query Match 2.4%; Score 33; DB 1; Le Best Local Similarity 84.6%; Pred. No. 0.75; Matches 33; Conservative 3; Mismatches 3; Conservative 3; Mismatches 3; Qy 1873 GAGATGGAGATGATGATGAAGATTGGGAAACACAAGAAT	Unknown. Unknown. Unclassified. 1 (bases 1 to 39) 1 (bases 1 to 39) Senoo, M., Watanabe, T. and Igarashi, K. Water-soluble mutein of FGF receptor, Patent: US 5750371-A 15 12-MAY-1998; Location/Qualifiers 1 39 /organism="unknown" 7 a 11 c 2 g 16 t	RESULT 1 AR007163/c AR007163 DEFINITION Sequence 15 from patent US 5750371. ACCESSION AR007163 VERSION AR007163 GI:3966647 KEYWORDS	12.4 12.4 12.4 12.4 15.4 12.4 15.4 15.4 15.4 16.4 17.4 17.4 18.4 18.4 19.5 19.4 19.5 19.4 19.5	326 12.8 0.9 17 1 327 12.4 0.9 14 1 329 12.4 0.9 14 1 330 12.4 0.9 14 1 331 12.4 0.9 15 1 332 12.4 0.9 15 1 333 12.4 0.9 15 1 334 12.4 0.9 15 1 336 12.4 0.9 15 1 337 12.4 0.9 15 1 338 12.4 0.9 15 1 339 12.4 0.9 15 1 330 12.4 0.9 15 1 331 12.4 0.9 15 1 331 12.4 0.9 15 1 332 12.4 0.9 15 1 333 12.4 0.9 15 1 334 12.4 0.9 15 1 336 12.4 0.9 15 1 337 12.4 0.9 15 1 338 12.4 0.9 15 1 339 12.4 0.9 15 1 340 12.4 0.9 15 1
linear PRI 09-MAY-2000 [3' region] [human, normal 1 of 2].	Length 39; 3; Indels 0; Gaps AT 1911	DNA and production thereof	linear PAT 04-DEC-1998	ACCESSION: AX598E ACCESSION: AX598E ACCESSION: AX6379 ACCESSION: E07289 ACCESSION: 177589 ACCESSION: AR0544 ACCESSION: AX1329 ACCESSION: AX3209 ACCESSION: AX3209 ACCESSION: AX708E	ACCESSION: 194532 ACCESSION: AR7105805 ACCESSION: AR714947 ACCESSION: BD123479 ACCESSION: A12791 ACCESSION: A40008 ACCESSION: A46520 ACCESSION: A76520 ACCESSION: AR7128998 ACCESSION: AR71026992 ACCESSION: AR7102669 ACCESSION: AR710266970 ACCESSION: AR7156870

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1 (bases 1 to 36)
Champion-Arnaud, P., Ronsin, C., Gilbert, E., Gesnel, M.C.,
Houssaint, E. and Breathnach, R.
Multiple mRNAs code for proteins related to the BEK fibroblast
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TK25=fibroblast growth factor receptor {3' region} [human, leukcyte DNA, Genomic, 36 nt, segment 1 of 2].
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GenBank staff at the National Library of Medicine created this entry [NCBI gibbsq 41355] from the original journal article. This sequence comes from Fig. 4b.
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Oligonucleotide OAB984
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Yayon, A., Ornitz, D.M., Klagsbrun, M., Leder, P. and Flanagan, J.
System for assaying binding to a heparin-binding growth factor
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A29208.1 GI:1248929
Patent: US 5789182-A 5 04-AUG-1998;
Location/Qualifiers
                                                                        Unclassified
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Location/Qualifiers
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Senoo, M., Watanabe, T. and Igarashi, K.
Water-soluble muttein of FGF receptor,
Patent: US 5750371-A 16 12-MAY-1998;
                                                                                                                                                                                                                                         AR007164
Sequence 16 from
AR007164
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Keating,M.T. and Morris,C.A.
Diagnosis of Williams syndrome and Williams
profile by analysis of the presence or abser
Patent: US 5858662-A 4 12-JAN-1999;
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I32953
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Wilson, S.E.

Methods and treatments for corneal healing
Patent: US 5703047-A 13 30-DEC-1997;
Patent: Location/Qualifiers
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Jun,S., Eriko,T., Chika,H., Akihiro,I., Masahiro,T.
Immortalized human papilla pili cell and method for
growth stimulants with the use of the same
Patent: JP 1999089565-A 22 06-APR-1999;
SHISEIDO CO LTD
OS Unidentific
PN JP 19990899
                                                                                                                                      E23733.1 GI:13024481
JP 1999089565-A/22.
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Patent: US 5589451-A 14 31-DEC-1996;
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C12R1:91),
PC (C12P21/02,C
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Sequence 13 from Patent W09705278.
A59560
Andersson,L., Moller,M.J., Wales,R., Siggens,K.W. and Plastow,G. METHODS FOR DETERMINING THE COAT COLOUR GENOTYPE OF A PIG Patent: WO 9705278-A 13 13-FEB-1997;
DALGETY PLC (GB)
                                                                                 unidentified unidentified
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Senoo, M., Watanabe, T. and Igarashi, K.
Water-soluble mutein of FGF receptor,
Patent: US 5750371-A 17 12-MAY-1998;
Location/Qualifiers
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                                                                                                                      A59560.1 GI:3714872
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/mol_type="genomic DNA"
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1 (bases 1 to 22)
Plastow,G.S. and Wales,R.
PETHODS FOR ANALYZING ANIMAL PRODUCTS
PATENT: WO 9854360-A 24 03-DEC-1998;
PLASTOW GRAHAM STUART (GB); WALES RICHA
LOCATION/Qualifiers
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Sequence
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AR128123.
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A82436
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20; Conserv
                                                                                                           Methods for determining the coat color Patent: US 6183955-A 13 06-FEB-2001;
                                                                                                                               Andersson, L., Moller, M. Johansson., Wales, R., Plastow, G. Stuart.
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/mol_type="genomic_DNA"
/db_xref="taxon:32644"
                                                                       /organism="unknown"
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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artificial sequences.
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Sequence 42 from Patent
AX203707
                                                                                               Methods and compositions for antisense vegf oligonucleotides Patent: WO 0152904-A 42 26-JUL-2001;
                                                                                                                         Gill, P.S. and Masood, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PF 27-MAY-1998 JP 199500368
PR 30-MAY-1997 GB 9711214.8,31-JAN-1998 GB
LBIF ANDERSSON, JAMES KIJAS, ELISABETTA GIUFFRA, GARY
EVANS, RICHARD WALES,
PI GRAHAM STUART PLASTOW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Methods for analyzing animal products Patent: JP 2002504814-A 24 12-FEB-2002; PIG IMPROVEMENT CO UK LTD
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12-FEB-2002
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                                                                                       Parkash, S. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism='Unidentified'
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
              note="VEGFR-1
                        /mol_type="genomic DNA"
/db_xref="taxon:32630"
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                                                 organism="synthetic construct"
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 AR019576
Sequence
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Sakurada,K., Palmer,T. and Gage,F.H.
Sakurada,K., Palmer,T. and Gage,F.H.
Regulation of tyrosine hydroxylase expression
Patent: US 6312949-A 39 06-NOV-2001;
Location/Qualifiers
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AR177699
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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from patent US 5783666.
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             PAT 05-DEC-1998
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APC (adenomatons malaria)
             1 (bases 1 to 24)
Albertsen,H., Anand,R., Carlson,M., Groden,J., Hedge,P.J.,
Joslyn,G., Kinzler,K., Markham,A.F., Nakamura,Y., Thliveris,A.,
Vogelstein,B. and White,R.L.
Diagnosis for mutant APC by immunoassay
Patent: US 6413727-A 61 02-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 24)
Albertsen, H., Anand, R., Carlson, M., Grod
Joslyn, G., Kinzler, K., Markham, A. Fred.,
Vogelstein, B. and White, R. L.
Detection of APC proteins
Patent: US 6114124-A 61 05-SEP-2000;
                                                                                                         Unknown
                                                                                                                                                          Sequence 61 from patent US
AR217050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 61 from patent US AR109537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APC (adenomatous polyosis coli) protein Patent: US 5783666-A 61 21-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unknown
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3 c 4 g
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Pred. No. 56;
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ed., Nakamura,Y., Thliveri
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155708/c
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176485/c
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155708
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1 (bases 1 to 24)

1 (bases 1 to 24)

Albertsen, H., Anand, R., Carlson, M., Grod Joslyn, G., Kinzler, K., Markham, A. Fred.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 24)
Albertsen, H., Anand, R., Carlson, M., Groden, J.
Joslyn, G., Kinnler, K., Markham, A., Nakamura, Yogelstein, B. and White, R.L.
Detection of inherited and somatic mutations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      colorectal cancer of humans
Patent: US 5648212-A 61 15-JUL-1997;
Location/Qualifiers
                                                                                                                                                                                                                    Unknown
                                                                                                                                                                                                                                Unknown.
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                                                                                                                                                            Vogelstein, B. and White, R.L.
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                                                                                        /organism="unknown"
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AX129008
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ORGANISM
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AX020546/c
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JOURNAL
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Sequence 226 :
AX129008
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Homo
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                                                                                                                                                                                                                                                                                  A method for identifying and characterizing Patent: WO 9934016-A 272 08-JUL-1999; GENENA LTD (IL); VIDER BEN ZION (IL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A method for identifying and characterizing Patent: WO 9934016-A 46 08-UUL-1999; GENENA LTD (IL); VIDER BEN ZION (IL) Location/Qualifiers
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Sequence 46 from Patent WO9934016.
AX020546
                       AX129008.1
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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sapiens (human)
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/mol_type="genomic DNA"

/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Primates;
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Pred. No. 53;
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AR116481/c
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                                                                                     Sequence 62
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AR116481.1
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AX129009
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18; Conserv
1 (bases 1 to 20) McKay,R., Dean,N., Monia,B.P., Nero,P.S. and Gaarde,W.A. Antisense oligonucleotide compositions and methods for t
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Ribozyme therapy for the treatment of proliferative
                                      Unclassified.
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                                                                                                                                                                                                                                                                                                                                                                                  Patent: WO 0130362-A 227 03-MAY-2001;
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/db xref="taxon:9606"
/note="Cdk2 ribozyme binding
5 c 5 g 4 t
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/mol_type="genomic DNA"
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                                                       Patent: JP 2001514905-A 62 18-SEP-2001;
ISIS PHARMACEUTICALS INC
OS Artificial Sequence
PN JP 2001514905-A/62
DD 18-SEP-2001
PF 07-AUG-1998 JP 2000509875
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              NERO, WILLIAM A GAARDE
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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C12Q1/68, A61K31/7088, A61K48/00, A61P35/00, C12N15/09, C12P19/34,
                    07-AUG-1998 JP 2000509875
13-AUG-1997 US 08/910629
ROBERT MCKAY,NICHOLAS DEAN,BRETT P MONIA,PAMELA SCOTT
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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AR211982
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Sakurada,K., Palmer,T. and Gage,F.H.
Regulation of tyrosine hydroxylase expression
Patent: US 6312949-A 40 06-NOV-2001;
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Ward,D.T. and Watt,A.T.
Antisense modulation of RECQL2 expression
Patent: US 6399378-A 38 04-JUN-2002;
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/mol_type="genomic DNA
/db_xref="taxon:32630"
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Pred. No. 6
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Pred. No. 68;
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Sequence 44
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C12R1:91),
PC (C12P21/02,
Strandedness: (
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                                                                              1 (bases 1 to 21)
Matson,R.S., Coassin,P.J., Rampal,J.B.
Oligonuclectide repeat arrays
Patent: US 5981185-A 44 09-NOV-1999;
                                                                                                                                    Unclassified
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MASAHIRO TAJIMA,
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JP 1999089565-A/24.
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06-APR-1999
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                             /organism="unknown"
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)_xref="taxon:32644"
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Pred. No. 68;
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Best Local Similarity 85.7%;
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Matson,R.S., Coassin,P.J., Rampal,J.l
Oligonucleotide repeat arrays
Patent: US 5981185-A 65 09-NOV-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                          Homo sapiens
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Unclassified.
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Mammalia; Eutheria; Primates;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 88;
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Pred. No. 89;
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AR139522/c
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1 (bases 1 to 20)
1 (bases 1 to 20)
1 (conrad, D.H. and Kelly, A.E.
1 (conrad, Conrad, A.E.
2 (conrad, Conrad, Conra
                                                                                                                          1 (bases 1 to 20)
Monia, B.P., Gaarde, W.A., Freier, S.M. and
Antisense modulation of TERT expression
Patent: US 6492171-A 61 10-DEC-2002;
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AR139522
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Keating,M.T. and Splawski,I.

Mutations in and genomic structure of HERG--a long

Patent: US 6207383-A 39.27-MAR-2001;
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AR079558
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Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or crelated to levels of vascular endothelial growth fact
Patent: US 6346398-A 7597 12-FEB-2002;
  Sequence
            AX482165
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Patent: WO 02086113-A 25 31-OCT-2002;
Isis Innovation Limited (GB)
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van der Kuyl, A.C. and Cornelissen, M.
Means and methods for treatment evaluation
                                                                                               AX721765
Sequence 144 from Patent
AX721765
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AUTHORS
TITLE
                                                                      JOURNAL
FEATURES
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ORGANISM
                                                                                                                                                                                                                                RESULT 47
AR222435/c
LOCUS
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AUTHORS
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ACCESSION
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ACCESSION
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A95063/c
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A95063
                                                                    Nucleotide sequences encoding the tomato light hypersensitive phenotype, encoded proteins and uses thereof Patent: US 6429299-A 5 06-AUG-2002;
Location/Qualifiers
                                                                                                                     1 (bases 1 to 19)
Bowler, C. and Mustilli, A.C.
                                                                                                                                                          Unknown
                                                                                                                                                                      Unknown.
                                                                                                                                                                                              AR222435.1
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BOWLER CHRIS (GB); STAZIONE ZOOL ANTON DOHRN (IT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 19)
Bowler, C. and Mustilli, A.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      unidentified
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                                                                                                                                               Unclassified.
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                                   /organism="unknown"
6 c 3 g
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/db xref="taxon:32630"
/note="primer 3'TAG019GENE-2"
8 c 3 g 5 t
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Pred. No. 1e+02;
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RESULT 48
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AR103195
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Bowler,C. and Mustilli,A.C.
Nucleotide sequence encoding tomato photosensitive
protein encoded thereby and utilization thereof
Patent: JP 2001526034-A 4 18-DEC-2001;
STAZIONE ZOOLOGICA ANTON DOHRN
                                                                  1 (bases 1 to 20)
Yuan, J. and Miura, M.
Programmed cell death genes and proteins
Patent: US 6087160-A 89 11-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleotide sequence encoding tomato protein encoded thereby and utilizar
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C12N15/09,A01H5/00,C07K14/415,C12N15/00
Nucleotide sequence encoding tomato photosensitive
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Key Location/Qualifiers
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JP 2001526034-A/4
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09-DEC-1997 IT RM97A000760
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                       organism="unknown"
8 c 1 g
                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                             /mol_type="genomic DNA"
/db_xref="taxon:32630"
6 c 3 g
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AX147705
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AX298917
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Mus sp.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Mus
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AX298917
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Clostridium perfringens
Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae;
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Sequence 7 from Patent WO0136673.
AX147705
                                                                                                                                                                                                            Bachmanov, A.A., Beauchamp, G.K., Chatterjee, A
Li, X., Ohmen, J.D., Reed, D.R., Ross, D. and To
Gene and sequence variation associated with
compounds and other sweeteners
Patent: WO 0183749-A 551 08-NOV-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Test for micro-organisms
Patent: WO 0136673-A 7 25-MAY-2001;
Creatogen Aktiengesellschaft (DE)
Location/Qualifiers
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                                                                                                                                                                                                  WARNER-LAMBERT COMPANY (US) ;
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GCTCAAGGAAGGACACA 17
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/mol_type="genomic DNA"
/db_xref="taxon:10095"
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/mol_type="genomic DNA"
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Pred. No. 1.1e+02;
0; Mismatches 1,
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Pred. No. 1.1e+02;
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YAKULT HONSHA CO LTD, YAKULT BIOSCIENCE KENKYU ZAIDAN
OS Artificial Sequence
PN JP 2001112485-A/11
PD 24-APR-2001
PF 19-OCT-1999 JP 1999296815
PI KIKUJI ITO, EISAKU KIKUCHI, TAKAHIRO MATSUKI, YUKIKO MIYAMOTO
C12N15/09,C12Q1/68,C12N15/00
CC
FH Key Location/Qualifiers.
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Patent: US 5827726-A 3 27-OCT-1998;
Docation/Qualifiers
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/db_xref="taxon:32630" 5 t
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Sequence 24
AR268232
  Vider, B.Z.

A method for identifying and characterizing Patent: WO 9934016-A 269 08-JUL-1999; GENENA LTD (IL); VIDER BEN ZION (IL)
                                                                                                                                                                                                                                                                                                                                                                                                                  Unknown.
Unknown.
                                                               Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                               Griffais,R., Hoiseth,S.K., Zagursky,R.J.,
Sankaran,B. and Fletcher,L.D.
Chlamydia pneumoniae polymucleotides and
Patent: US 6559294-A 5279 06-MAY-2003;
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A method for identifying and characterizing Patent: WO 9934016-A 275 08-JUL-1999;
GENENA LTD (IL); VIDER BEN ZION (IL)
                                                                                                                                          Novel human gene relating to respiratory diseases, obesity, and inflammatory bowel disease Patent: WO 0178894. 140 25-OCT-2001; Genome Therapeutics Corp. (US)
                                                                                                                                                                                                                                     synthetic
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                                                                                                                                                                                                                                                 synthetic construct
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Mammalia; Eutheria; Primates;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                           organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:9606"
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                                                               note="Primer"
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PATENT: JP 1996308586-A 2 2.
CHUGAI PHARMACEUT CO LTD
OS None
OC Artificial sequences.
PN 196308586-A/2
PD 26-NOV-1996
PF 15-MAR-1996 JP 19960874
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17; Conserv
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                                                                                                                                                                                                                                                                                          E12301
E12301.1 GI:3251135
JP 1996308586-A/2.
strandedness: Single;
CC topology: Linear;
CC hypothetical: No;
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JP 1996308586-A/2
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ilarity 85.0%;
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                                                                                                                                                                                                                                                                                corresponding mutant line and a link to a database providing a graphical display of the insertion site are available at http://dbsgap.versailles.inra.fr/publichnes/. This sequence he been generated in the framework of the French plant genomics program 'Genoplante' (http://www.genoplante.com and http://genoplante.info.infobiogen.fr).
                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (21-NOV-2002) Balzergue S., UMRGV, INRA/CNRS, 2 rue Gaston Cremieux, 91057 Evry cedex, FRANCE
PCR was performed on DNA from transformants of Arabidopsis thaliana plants from INRA (Versailles). The DNA fragment(s) resulting from the PCR were directly sequenced from the left or the right border to determine the genomic sequence flanking the insertion. T-DNA derived sequences were removed. Information to order the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Brunaud, V., Balzergue, S., Dubreucq, B., Aubourg, S., Samson, F., Chauvin, S., Bechtold, N., Cruaud, C., DeRose, R., Pelletier, G., Lepiniec, L., Caboche, M. and Lecharny, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AJ527748.1 GI:26796008
left border, T-DNA flanking sequence.
Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
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Location/Qualifiers
                                                                                                                 /clone_lib="Arabidopsis thaliana
                                                                                             /note="T-DNA flanking
                                                                                                                                                                                         /mol_type="genomic DNA"
/cultivar="Wassillewskija"
                                                                                                                                                                                                                              organism="Arabidopsis thaliana"
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                                                                                                                                                     clone="146F12"
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ATH527822/c
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Best Local
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                                                                                                                                                                                                                                     l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR was performed on DNA from transformants of Arabidopsis thaliana plants from INRA (Versailles). The DNA fragment(s) resulting from the PCR were directly sequenced from the left or the right border to determine the genomic sequence flanking the insertion. T-DNA derived sequences were removed. Information to order the corresponding mutant line and a link to a database providing a graphical display of the insertion site are available at http://dbsgap.versailles.inra.fr/publiclines/. This sequence has
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17;
tagged site.
L77371
                                  DOGHOX7B 20 bp
Canis familiaris Homeobox 7 (HOX7)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    been generated in the framework of the French plant genomics program 'Genoplante' (http://www.genoplante.com and http://genoplante-info.infobiogen.fr)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (21-NOV-2002) Balzergue S., UMRGV, INRA/CNRS, 2 rue Gaston Cremieux, 91057 Evry cedex, FRANCE
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left border; T-DNA flanking sequence.
Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of pre-insertion sites
EMBO Rep. 3 (12), 1152-1157 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Brunaud, V., Balzergue, S., Dubreucq, B., Aubourg, S., Samson, F., Chauvin, S., Bechtold, N., Cruaud, C., DeRose, R., Pelletier, G., Lepiniec, L., Caboche, M. and Lecharny, A. T.-DNA integration into the Arabidopsis genome depends on sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Arabidopsis thaliana T-DNA flanking sequence, left border, clone
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                                                                                                                                                                                                                                                                                                                                                     /note="T-DNA flanking sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Arabidopsis thaliana"
/mol_type="genomic DNA"
/cultivar="Wassillewskija"
                                                                                                                                                                                                                                                                                                                                                                                            clone_lib="Arabidopsis thaliana
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DOGKIT1A
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Best Local Similarity
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Matches 17; Conserv
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primer_bind
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Venta, P.J., Brouillette, J.A., Yuzbasiyan-Gurkan, V. and Brewer, G.Gene-specific universal mammalian sequence-tagged sites:
application to the canine genome
Unpublished (1996)
Original source text: Canis familiaris DNA.
Gene-specific universal mammalian sequence-tagged site for KIT1.
Primer for the 5' end is in exon 18. Human product is 650 bp.
Canine product is 650 bp. PCR conditions: 1 min, 94 C, 2 min, 57 (3 min, 72 C, 35 cycles.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Original source text: Canis familiaris DNA.

Gene-specific universal mammalian sequence-tagged site for HOX7.

Primer for the 3' end is in exon 2. Human product is 151 bp. Canine product is 151 bp. PCR conditions: 1 min, 94 C, 2 min, 57 C, 3 min, 72 C, 35 cycles.
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STS; Home
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Canis familiaris c-KIT Protooncogene (KIT1)
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Venta, P.J., Brouillette, J.A., Yuzbasiyan-Gurkan, V.
Gene-specific universal mammalian sequence-tagged
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                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Canis familiaris
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Mammalia; Eutheria; Carnivora;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Canis familiaris (dog)
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                                                                                      complement(1. .20)
/note="PCR primer binding site"
                                                                                                                                                                                                                    Location/Qualifiers
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/mol type="genomic DNA"
/db_xref="taxon:9615"
1. .20
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/mol_type="genomic DNA"
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5 c 7 g
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Pred. No. 1.1e+02;
 Score 15.2; DB 1;
Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             n; PCR primer; mammalian STS
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; Fissipedia; Canidae;
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                Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                        Euteleostomi; Canis.
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KEYWORDS
SOURCE
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AR133380
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AR133382
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DEFINITION
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Matches 15; Conserv
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Unclassified
                   Unknown.
                                       AR133382.1
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Stinchcomb,D.T., Jarvis,T. and McSwigg
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 1806 27-FEB-2001;
                                                                                                                                                                                                                                                                                           Unknown
                                                                                                                                                                                                                                                                                                                         AR133381.1
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Stinchcomb, D.T., Jarvis, T. and McSwiggen, J.
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Sequence 42 from patent US
AR106794
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AX553634
AX553634.1 GI:25897632
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                                                                                Antisense inhibition of G-alpha-16 expression Patent: US 6107091-A 42 22-AUG-2000; Location/Qualifiers
                                                                                                                    1 (bases 1 to 18)
Cowsert, L.M.
                                                                                                                                             Unclassified.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
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Mammalia; Eutheria; Primates;
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Bennett, C.Frank. and Cowsert, L.M.
Antisense modulation of Her-4 expression
Patent: US 6255111-A 65 03-JUL-2001;
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/mol type="genomic DNA"
/db_xrefs"taxon:9606"
/note="Cdc25 hs ribozyme binding
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       1 (bases 1 to 17)
Donahoe, P.K., Gustafson, M. and He, W.W.
DNA encoding a receptor for Mullerian inhibitory substance, and corresponding vectors, cells, probes, and recombinant me Patent: US 5547854-A 10 20-AUG-1996;
                                                                                                                                          Sequence 10 from patent I25016
                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 17)
Donahoe, P.K., Gustafson, M., He, W.-W. and
Nucleic acids encoding a TGF.-beta. type
Patent: US 5538892-A 10 23-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 10 from patent I23857
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unknown
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Homo sapiens
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Mammalia; Eutheria; Primates;
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                                                                          Unclassified.
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Similarity 88.9%;
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nilarity 82.4%;
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2 c 4 g
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/mol_type="genomic DNA"
/db xref="taxon:9606"
/noTe="Cdc25 hs ribozyme binding
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Pred. No. 1.2e+02;
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1 receptor
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                                                                                                                                                                         Robbins, J.M. and Tritz, R.
Ribozyme therapy for the treatment of
                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/organism="unknown"
a 2 c 4 g
                                                 /note="IL-1 beta ribozyme 3 c 5 g 6
                                                                        /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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3 c
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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            93.8%;
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            Score 14.4; DB 1
Pred. No. 1.1e+02
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Pred. No. 1.2e+02;
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AR048076
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synthetic construct
artificial sequences.

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1 (bases 1 to 16)
1 retarding the conversion of hormone-dependent cancer into hormone-independent cancer
hormone-independent cancer
Patent: JP 2000178202-A 6 27-JUN-2000;
TAKEDA CHEMICAL INDUSTRIES LTD
OS ARTIFICIAL Sequence
PN JP 2000178202-A/6
                                                                                                                                                                                                   AR048076
Sequence 17 from patent
AR048076
             15; Conserv
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Karn, ..., Gait, M.John., Heaphy, S. and Dingwall, C.
RNA oligonuclectides that bind HIV tat protein
Patent: US 5821046-A 17 13-OCT-1998;
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BD002055.1 GI:18628795
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Similarity 75.0%;
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A61K38/04, A61K38/22, A61K45/00, A61P13/08, A61P35/00//C07K7/23
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Location/Qualifiers
                                                      /organism="unknown"
4 c 5 g
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/mol_type="genomic DNA"
/db_xref="taxon:32630" 2 t
5 c 1 g 2 t
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Pred. No. 1.2e+02;
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Pred. No. 1.1e+02;
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Unclassified.
Karn, J., Gait, M.J., Heaphy, S. and Dingwall, C.
          Unclassified.

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1 (bases 1 to 17)

1 (bases 1 to 17)

Karn,J., Gait,M.John., Heaphy,S. and Dingwall,C. RNA oligonucleotides that bind HIV tat protein Patent: US 5821046-A 20 13-CCT-1998;

1. .17
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Karn, J., Gait, M.J., Heaphy, S. and Dingwall, C. Viral (HIV) growth inhibition
Patent: US 6114109-A 17 05-SEP-2000;
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Pred. No. 1.2e
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Patent: US 6114109-A 20 05-SEP-2000;
Location/Qualifiers
1. 17
                                                                                                                Narn, J., Gait, M. John., Heaphy, S. and Dingwall, C. RNA oligonucleotides that bind HIV tat protein Patent: US 5821046-A 23 13-CCT-1998; Location/Qualifiers
                                                                                                                                                                                                                                   AR048082
Sequence 23 from patent US
AR048082
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Pavco,P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398-A 5715 12-FEB-2002;
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AR108985
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AR220012
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Cowsert, L. M.
Antisense modulation of G-alpha-11 expression
Patent: US 5951455-A 30 14-SEP-1999;
          1 (bases 1 to 18)
Anderson, S. and Banta, S.
Design and production of mutant 2,5-diketo-D-gluconic
reductase enzymes with altered cofactor dependency
Patent: US 6423518-A 5_23-JUL-2002;
                                                                          Unknown
                                                                                                                                                                                                                                                                                                                              Unclassified.

(bases 1 to 18)

Karn,J., Gait,M.J., Heaphy,S. and Diviral (HIV) growth inhibition

Patent: US 6114109-A 23 05-SEP-2000;
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Patent: WO 0157206-A 1035 09-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US) ; Fattaey,
Location/Qualifiers
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Sequence 1035 from Patent
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Patent: WO 0157206-A 1034 09-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US) ; Fattaey, Ali R.
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/mol_type="mRNA"
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/mol_type="mRNA"
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/mol type="mRNA"
/db_xref="taxon:32630"
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Sequence 28 from Patent WO9925815.
A95626
                                             Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J. Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398_A 4205 12-FEB-2002;
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Herrmann, B. and Kispert, A.
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HERRMANN BERNHARD (DE); MAX PLANCK GESELLSCHAFT
Location/Qualifiers
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WO 9202228-A 17 20-FEB-1992;
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/organism="unknown" 5 g
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol_type="mRNA"
/db_xref="taxon:32630"
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Pavco, P., McSwiggen, J.; Stinchcomb, D. and Escobedo, J. Method and reagent for the treatment of diseases or crelated to levels of vascular endothelial growth fact Patent: US 6346398-A 4206 12-FEB-2002;
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Pavco,P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 4242 12-FEB-2002;
Location/Qualifiers
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                                          1 (bases 1 to 17)
Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J. Method and reagent for the treatment of diseases or related to levels of vascular endothelial growth fact Patent: US 6346398_A 5779 12-FEB-2002;
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PavCo,P., McSwiggen,J., Stinchcomb,D. and Escober Method and reagent for the treatment of diseases related to levels of vascular endothelial growth Patent: US 6346398-A 5714 12-FEB-2002;
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Pavco,P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 4357 12-FEB-2002;
                                                                                                          Unclassified
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Pred. No. 1.4e+02;
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AR190292
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Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 5817 12-FEB-2002;
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Pavco,P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 5780 12-FEB-2002;
Location/Qualifiers
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                                                           1 (bases 1 to 17)
Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 7685 12-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                 ravco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 7684 12-FEB-2002;
Location/Qualifiers
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Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 7674 12-FEB-2002;
Docation/Qualifiers
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                       /organism="unknown"
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Pred. No. 1.4e+02;
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Pred. No. 1.4e+02;
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AUTHORS
TITLE
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AR192199
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AR286013/c
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1 (bases 1 to 17)

Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor related to 1846398-A 7686 12-FEB-2002;

Patent: US 6346398-A 7686 12-FEB-2002;

1 1 17----"unknown"
         AR286013
Sequence 385
AR286013
AR286013.1
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AR192199
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15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
15; Conserv
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           GI:29723609
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Pred. No. 1.4e.
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AX008727
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AR286406
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Sequence 1 from Patent WO9964625.
AX008727
                                                                                                                                               Human immunodeficiency virus
Human immunodeficiency virus
Viruses; Retroid viruses; Retroviridae;
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15; Conserv
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Beigelman, L., Burgin, A.,
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Beigelman, L., Burgin, A., Beaudry, A., Karpeisky, A.,
Matulic-Adamic, J., Sweedler, D. and Zinnen, S.
Synthetic ribonucleic acids with RNAse activity
Patent: US 6528640-A 385 04-MAR-2003;
1. 17
                                                          Methods and kits for discovery of rna-binding
Patent: WO 9964625-A 1 16-DEC-1999;
RIBOTARGETS LIMITED (GB)
Location/Qualifiers
                                                                                                            Prescott, C.D. and Karn, J.
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3 c 6 g
/mol_type="genomic RNA
/db_xref="taxon:12721"
4 c 5 g
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                                     organism="Human immunodeficiency
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Pred. No. 1.4e+02;
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                                                                                                                                                Lentivirus; Primate
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VERSION
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Patent: WO 0159103-A 3622 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Bla
McSwiggen, James (US); Chowrira, Bharat
Location/Qualifiers
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AX215402
AX215402.1
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synthetic construct
artificial sequences.
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15; Conserv
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AX218180
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/mol type="mRNA"
/db_xref="taxon:32630"
/note="Nucleic Acid"
8 c 7 g 1 t
                                                      /organism="synthetic construct"
/mol_type="mRNA"
/db xref="texon:32630"
/note="Nucleic Acid"
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AX218315
RESULT 113
AX325661
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Patent: WO 0159103-A 3757 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
MCSwiggen, James (US); Chowrira, Bharat M. (
Location/Qualifiers
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Sequence 430 from Patent
AX227058
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/mol type="mRNA"
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/mol_type="mRNA"
/db_xref="taxon:32630"
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Pred. No. 1.4e+02;
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l Similarity 88.2%;
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AX325662
AX325662.1 GI:18096421
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Sequence 1799 from Patent WO0192512.
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AX498838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Solanum tuberosum Solanum tuberosum
          AX498838.1 GI:23381120
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                                                                                                                                                                                                                                                                                                                                          Kmiec,E.B., Gamper,H.B., Rice,M.C. and Kim,J. Targeted chromosomal genomic alterations in p single stranded oligonucleotides Patent: WO 0192512-A 1800 06-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Solanum tuberosum (potato)
Solanum tuberosum
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Spermatophyta; Magnoliophyta; eudicotyledons; core e
Asteridae; lamiids; Solanales; Solanaceae; Solanum.
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/mol_type="genomic DNA"
/db_xref="taxoni4113"
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/mol type="genomic DNA"
/db_xref="taxon:4113"
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AX500369/c
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Human posh-like protein 1
Patent: EP 1239051-A 122 11-SEP-2002;
Aeomica, Inc. (US)
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Sequence 122 from Patent
AX530613
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                                   Shannon, M.
                                                        Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens (human)
                                                                            Homo sapiens (human)
Homo sapiens
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Similarity 88.2%;
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                                                                                                                                                                                                                                                                                                                                                                                           Zhan,J.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
2 c 1 g
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 1.4e+02;
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Pred. No. 1.4e+02;
0; Mismatches 2;
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AX673357
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AX530614/c
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Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                              Patent: WO 03004526-A 1802 16-JAN-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                        medicines
                                                                                                                                Telerman, A., Amson, R. and Tuijnder, M. Sequences involved in phenomena of tumour
                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Patent: EP 1239051-A 123 11-SEP-2002;
Aeomica, Inc. (US)
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Sequence 123 from Patent
AX530614
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/organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
6 c 2 g
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/mol_type="genomic DNA"
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/db_xref="taxon:9606"
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Matches 15; Conserv
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2532 GGTAGAAGACTTGGATC 2548
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15; Conservative
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                                                                                                                                                                                                                                                                                                     Mus musculus
                                                                                                                                                                                                                                                                                                                                                            Sequence 3780 from Patent AX726093
                                                                                                                                                                 Patent: WO 03025176-A 3780 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                                                                                                                                    Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression, tumor reversion, apoptosis and/or virus resistance and their use
                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
/db_xref="taxon:10090"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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                                          Score 13.8; DB 1;
Pred. No. 1.4e+02;
0; Mismatches 2;
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Pred. No. 1.4e+C
O; Mismatches
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Pred. No. 1.4e+02;
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                                                                                                                                                                                                                                        Telerman, A., Amson, R. and Tuijnder, M.
Sequences involved in phenomena of tumour suppression, tumo
                                                                                                                                                                                                                                                                                                                                                                                   AX733343
Sequence 4977 from Patent
AX733343
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                                                                                                                                                                                                                                                                                                                              Homo sapiens (human)
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as
                                                                                                                                                                                                 Molecular Engines Laboratories (FR)
                                                                                                                                                                                                                              medicines
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Molecular Engines Laboratories (FR)
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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PF 18-NOV-1998 JP 2000521181

PF 18-NOV-1997 EP 97120190.0,02-MAR-1998 EP 98103596.7 PI

BERNHARD HERRMANN, BIRGIT KOSCHORZ, ANDREAS KISPERT PC

C12N15/09, A01K67/027, A61K31/7088, A61K38/45, A61K39/395, A61K48/ PC

00, A61P15/16,

C07K16/40,C12N1/15,C12N1/19,C12N1/21,C12N5/10 C12P21/08,C12N15/00 C12P21/08,C12N15/00 CCC

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MAX PLANCK GESELLSCHAFT ZUR FOERDERUNG DER WISSENSCHAFTEN EV
OS Artificial Sequence
PN JP 2001523449-A/19
PD 27-NOV-2001
PF 18-NOV-1998 JP 2000521181
PF 18-NOV-1997 EP 97120190.0,02-MAR-1998 EP 98103596.7
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1 (bases 1 to 17)
Herrmann,B., Koschorz,B.
Nucleic acids involved in
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Zhang, J.

A human G protein coupled receptor Patent: WO 03031621-A 493 17-APR-2003; Amersham Biosciences (SV) Corp. (US)
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/mol type="genomic DNA"
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Patent: WO 9529259-A 4 02-NOV-1995;
STICHTING CENTRAPL LAB (NL)
Other publication AU 2319495 951116.
Location/Qualifiers
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Oligoribonucleotide 18-mer.
A21030
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Sequence 4 from Patent WO9529259.
A46964
          A88187
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Voorberg, J.J., Van, M.J. and Mertens,
METHOD AND MEANS FOR DETECTING AND T
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synthetic construct
artificial sequences.
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335 from Patent W09833904
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
2 c 4 g
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/mol_type="mRNA"
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A88187.1
unclassified.
1 (Dases 1 to 18)
Min,J.W. and Fiers,W.
Min,J.W. and Fiers,W.
NEW IMMUNOPROTECTIVE INFLUENZA ANTIGEN AND ITS USE IN VACCINATION
Patent: WO 9907839-A 48 18-FEB-1999;
VLAAMS INTERUNIV INST BIOTECH (BE); MIN JOU WILLY (BE)
                                                                                                                                                                                                                                                                                                                                                                                                                    I (bases 1 to 18)
Brysch, W.D. and Schlingensiepen, K.D.
An antisense oligonucleotide preparation method Patent: EP 0856579-A 335 05-AUG-1998;
BIOGNOSTIK GES (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 18)
Brysch, W. and Schlingensiepen, K.
AN ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD
Patent: WO 9833904-A 335 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
                                                                                     unidentified
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Sequence 335 from Patent EP0856579.
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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AR036682
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Hall,F.L., Nimni,M.E., Tuan,T.-L., Wu,L. and Cheung,D.T.
Artificial skin prepared from coclagen matrix containing
transforming growth factor.beta having a collagen bind
Patent: US 5800811-A 21 01-SEP-1998;
Location/Qualifiers
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                                                                                                                   1 (bases 1 to 18)
Karn, J., Gait, M.John., Heaphy, S. and Dingwall, C.
RNA oligonucleotides that bind HIV tat protein
Patent: US 5821046-A 13 13-CCT-1998;
Location/Qualifiers
                                                                                                                                                                                     Unknown
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/mol_type="genomic_DNA"
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AR077364
1 (bases 1 to 18)

Voorberg, J.Jacobus., van Mourik, J.Aart. and Mertens, K.

Method and means for detecting and treating disorders in the blood

coagulation cascade

Patent: US 6083905-A 4 04-JUL-2000;
                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 18)
Monia, B.P. and Coweert, L.M.
Antisense inhibition of EGR-1 expression
Patent: US 6008048-A 17 28-DEC-1999;
Location/Qualifiers
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Griffiths, A., David., Williams, S. Cameron., Wate
Nissim, A., Winter, G. Paul., Johnson, K. Stuart.
                                                                                                                      Sequence 4 from AR102333
                                                                                                                                                                                                                                                                                                                                                                                                                                                AR096633
Sequence 17 from patent US
AR096633
                                                                        Unknown.
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                                                                                    Unknown.
                                                                                                           AR102333.1 GI:12813131
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Pred. No. 1.5e+C
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Pred. No. 1.5e+0; Mismatches
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KEYWORDS
SOURCE
          RESULT 138
AR117984
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ACCESSION
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AR108975
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TITLE
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Best Local Similarity
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Sequence
AR106793
                                                                                                                                                             1 (bases 1 to 18)
Karn,J., Gait,M.J., Heaphy,S. and Dingwall,C.
Viral (HIV) growth inhibition
Patent: US 6114109-A 13.5.5.
                                                                                                                                                                                                                 Unknown
                                                                                                                                                                                                                                                       Sequence 13
AR108975
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Cowsert, L.M.
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                                              1 (bases 1 to 18)
Griffiths,A.D., Williams,S.C., Waterhouse,P.M., Nissim,A.Winter,G.P., Johnson,K.S. and Smith,A.J.H.
Methods for producing members of specific binding pairs
Patent: US 6492160-A 79 10-DEC-2002;
                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 18)
Nimni,M.E., Hall,F.L., Wu,L., Han,B. and Shors,E.C.
Bone morphogenetic proteins and their use in bone g
Patent: US 6352972-A 21 05-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Johnson, K. Stuart., Winter, G. Paul., Griffiths, A. David., Smith, A. John. Hammond. and Waterhouse, P. Michael. Methods for producing members of specific binding pairs Patent: US 6140471-A 33 31-OCT-2000;
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AR117984
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Sequence 79 from patent US 6492160.
AR265427
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AR198571
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Sato,S., Higashikuni,N., Kudo,T. and Kondo,M.
DNAs encoding new fusion proteins and processes for preparing useful polypeptides through expression of the DNAs patent: US 6506595-A 9 14-JAN-2003;
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15; Conserv
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15; Conservat
Shimkets,R.A. and Fernandes,E.
Nucleic acids and secreted polypeptides
Patent: WO 0110902-A 40 15-FEB-2001;
Curagen Corporation (US)
                                                                                                                                 Sequence 40 from Patent AX084246
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Sequence 19 from Patent W00104358.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hepatitis B virus
Hepatitis B virus
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Methods and kits for discovery of rna-binding Patent: WO 9964625-A 3 16-DEC-1999;
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Human immunodeficiency virus
Viruses; Retroid viruses; Retroviridae;
                                                                                         synthetic construct
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                                                                artificial sequences.
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/mol_type="genomic RNA"
/db_xref="taxon:12721"
db_x 4 c 6 g 3 t
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/db_xref="taxon:10407"
2 c 5 g 7 t
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Sequence 2 from Patent WO02095071.
AX708786
                                                                                         Means and methods for identifying genes and proteins involved
the prevention and/or repair of a replication error
Patent: WO 02095071-A 2 28-NOV-2002;
Koninklijke Nederlandse Akademie van Wetenschappen (NL)
                                                                                                                                                                                     synthetic construct
synthetic construct
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synthetic construct
artificial sequences.
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/note="Primer RO3C_C"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
/note="PCR PRIMER"
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                                                                                                        Schlingensiepen, K.H. and Brysch, W.
An antisense oligonucleotide preparation method
An antisense oligonucleotide preparation
Patent: JP 2001511000-A 335 07-AUG-2001;
BIOGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH
OS UNKNOWN
PN JP 2001511000-A/335
PD 07-AUG-2001
PP 30-JAN-1998 JP 1998532533
PR 31-JAN-1997 EP 97101531.8
PR 31-JAN-1997 EP 97101531.8
PI KARL HERMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
PC C12N15/11, C07H21/04, A61K31/70
CAN antisense oligonucleotide preparation method FH Ke
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18 bp
Sequence 87 from Patent WO03018837.
AX713201
                                                                                                                                                                                                                                                                                                                          unidentified
                                                                                                                                                                                                                                                                                                                                                                                             BD065700 18 bp
An antisense oligonucleotide pre
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                                                                                                                                                                                                                                                                                                                                                                    BD065700.1 GI:22611303
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                                                                                                  Location/Qualifiers
                                                      /organism='Unknown'.
Location/Qualifiers
/mol_type="genomic DNA
/db_xref="taxon:32644"
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/db_xref="taxon:32630"
/note="Oligonukleotid"
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                            organism="unidentified"
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88.2%;
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Pred. No. 1.5e+02;
0; Mismatches 2;
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SM synthetic construct
artificial sequences.

E 1 (bases 1 to 18)

E 1 (bases 1 to 18)

Novel insulin/IGF/relaxin family polypeptide and its DNA
L Patent: WO 0181562-A 10 01-WOV-2001;
TAKEDA CHEMICAL INDUSTRIES LTD, YASUAKI ITO, NOBUHIRO SUZUKI,
KAZUNORI MISHI, HIDEKI KIZAWA, MASATAKA HARADA, KAZUHIRO OGI
OS Artificial Sequence
PN WO 0181562-A/10
PD 01-NOV-2001 WO 2001JP003399
PF 20-ARR-2001 WO 2001JP003399
PF 20-ARR-2001 WO 2001JP003399
                                                                                                                                                                                                                                                1653
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Kit and method |
BD104044
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PC C12N15/09,C07K14/65,C07K16/26,C12N1/15,C12N1/19,C12N1/21 PC
PC C12N15/09,C07K14/65,C07K16/26,C12N1/15,C12N1/19,C12N1/21 PC
PC C12P21/02,C12P21/08,A61K38/17,A61K38/28,A61K38/30,A61K39/395
PC A61R45/00,
PC A61R45/00,A61P3/00,A61P19/00,A61P9/00,A61P5/00,A61P25/00, PC
A61P3/00,A61P3/00,A61P19/00,G01N33/50,G01N33/15 CC Primer
PH Key Location/Qualifiers
PH Key 1. 18
PT Source 1/organism='Artificial Sequence'.
                                                                                                                                                                                                               N
                                synthetic construct
synthetic construct
                                                                BD104044.1 GI:22649618
WO 0192572-A/148.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PF 20-APR-2001 WO 2001JF003399
PR 21-APR-2000 JP 00P 126340,03-JUL-2000 JP 00P 205587
10-AUG-2000 JP 00P 247962,22-DEC-2000 JP 00P 395050 PI
YASUAKI ITO,NOBUHIRO SUZUKI,KAZUNORI NISHI,HIDEKI KIZAWA, PI
MASATAKA HARADA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BD103233.1 GI:2
WO 0181562-A/10.
                  artificial sequences
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(bases 1 to 18)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C12P21/02,C12P21/08,A61K38/17,A61K38/28,A61K38/30,A61K39/395,A61K45/00,
                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                          /organism='Artificial Sequence'
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                            organism="synthetic construct"
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                                                                                                                                                                                                                                                                                            Score 13.8;
Pred. No. 1.
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Pred. No. 1.5e+02;
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                                                                                                                                  linear
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                                                                                                                                  27-AUG-2002
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E39157/c
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PI SHOGO
PC C12Q1
CC Descr
FH Key
FT Sourc
                                                                                                                                                                                                                                                                                 artificial sequences.

1 (bases 1 to 18)

Seiji,S., Masahiko,H., Toshiyuki,K. and Masaaki,K.

Seiji,S., Masahiko,H., Toshiyuki,K. and Masaaki,K.

DNA encoding novel fused protein and process for producing useful protein mediating the expression thereof

Patent: JP 199341991-A 3 14-DEC-1999;

ITO HAM KK,TUZO UDAKA

OS Artificial Sequence

PN JP 199341991-A/3
                                                                                                                                                                                                                                                                                                                                                                                                                               synthetic construct
synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA encoding novel fused protein and process for protein mediating the expression thereof.
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Patent: WO 0192572-A 148 06-DEC-2001;
NISSHINBO INDUSTRIES INC. SYSTEM RESEARCH INC,HIDETOSHI INOKO, TI
KAGIYA, TATSUO ICHIHARA,YOSHIYUKI MATSUMURA,SHOGO MORIYA,MICHIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTACCCTGCCACAGGAA
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C12Q1/68,C12M1/00,C12N15/09,G01N33/53
Description of Artificial Sequence:capture
Key Location/Qualifiers
                                                                                                source
                                                                                                                                                                                        C07K14/655,
                                                                                                                                                                                                     SEIJI SATO, MASAHIKO HIGASHIKUJI, TOSHIYUKI KUDO, MASAAKI KONDO C12N15/09, C12N1/21, C12P21/02, C12P21/02//C07K14/605, C07K14/62,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUN-2001 WO 2001JP004662
01-JUN-2000 JP 00P 164798
HIDETOSHI INOKO,TAEKO KAGIYA,TATSUO ICHIHARA,YOSHIYUKI
                                                                                                                                            C12N15/00, (C12N15/00, C12R1:08)
                                                                                                                                                                        C07K19/00, (C12N15/09, C12R1:08), (C12N1/21, C12R1:08), (C12P21/02
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06-DEC-2001
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                                                            /organism='Artificial Location/Qualifiers
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Location/Qualifiers
/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GI:13019231
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/db_xref="taxon:32630"
3 c 6 g
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Pred. No. 1.5e+02;
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OS ARTÉFICIAI Sequence
PN JP 199934191-A/4
PD 14-DEC-1999
PF 30-MAR-1999 JP 1999081
PF SELJI SATO, MASAHIKO H.
PC C12N15/09, C12N1/21, C1.
PC C07K14/655,
PC C12R1:08),
PC C12R1:08],
PC C12R
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Unclassified.

1 (bases 1 to 18)

Somenberg,N., Katze,M.G., Roy,S., Ko Anti-tumor agent assay using PKR Patent: US 5670330-A 19 23-SEP-1997;
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1 (bases 1 to 18)
Seiji,S., Masahiko,H.,
                                                                                                                               Unknown
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nilarity 88.2%;
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/db_xref="taxon:32630"
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Location/Qualifiers
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                                                  Koromilas, A.E. and Barber, G.H.
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15; Conserv
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195705
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Johnson, K., Stuart., Winter, G.Paul., Griffiths, A.David.,
Smith, A.John. Hammond. and Waterhouse, P. Michael.
Methods for producing members of specific binding pair;
Patent: US 5733743-A 33 31-MAR-1998;
                                                                                                                                                                       1 (bases 1 to 15)
Vogelstein,B., Kinzler,K.W., Zhang,L. and Gene expression profiles in normal and car Patent: US 6333152-A 233 25-DEC-2001;
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Sequence 233 from patent
AR180165
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                                                                                                                                                                                 and Zhou, W. cancer cells
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Local Similarity 93.3%;
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                                                                       BD065687 16
An antisense oligonucleotide
BD065687 GI:22611290
JP 2001511000-A/322.
                                                                                                                                                                                                                                  14;
1 (bases 1 to 16)
Schlingensiepen, K.H. and Brysch, W.
An antisense oligonucleotide preparation method
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Brysch, W. and Schlingensiepen, K.
ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD
PATENT: WO 9833904-A 322 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
                                      unclassified.
                                                  unidentified
                                                             unidentified
                                                                                                                                                                                                                                                                                                                                                       BIOGNOSTIK GES (DE)
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Brysch, W.D. and Schlingensiepen, K.D.
An antisense oligonucleotide preparation
Patent: EP 0856579-A 322 05-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                              unidentified
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
7 c 2 g
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/db_xref="taxon:32644"
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Pred. No. 1.5e+02;
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Other
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BIOGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH
OS UNKNOWN
PN JP 2001511000-A/322
PD 07-AUG-2001
PF 30-JAN-1998 JP 1998532533
PR 31-JAN-1997 EP 97101531.8
PR 31-JAN-1997 EP 97101531.8
PI KARL HERMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
PC C12N15/11,C07H21/04,A61K31/70
CC An antisense oligonucleotide preparation method FH
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Sequence 10 from Patent WO9604387.
A48870
                                                   AR085293
Sequence 5 from patent
AR085293
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         unidentified
                                      AR085293.1
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Similarity 93.3%;
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larity 93.3%;
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publication FR 2723378 960209
Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Location/Qualifiers
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/db_xref="taxon:32644"
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                                         GI:10012062
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Pred. No. 1.6e+02;
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AR127158
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Kornbluth, J.
Natural killer lytic a
Patent: US 5981705-A 5
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Sequence
AR188867
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14; Conserv
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AR127158
                                                                                   1 (bases 1 to 17)
Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or corelated to levels of vascular endothelial growth facts
Patent: US 6346398-A 4355 12-FEB-2002;
                                                                                                                                                   Unknown
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DNA sequences coding for the human proteins the interleukin-lbeta converting enzyme Patent: US 6180386-A 10 30-JAN-Z001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unknown.
Unclassified.
1 (bases 1 to 17)
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                                                                                                                                          Unclassified.
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Score 13.4; D
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Pred. No. 1.6e+02;
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                      DB 1;
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Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398.A 4356 12-FEB-2002;
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                                                          Sequence 2375 from Patent AX216933
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Beigelman, L., Burgin, A., Beaudry, A., Karpeisky, A.,
Matulic-Adamic, J., Sweedler, D. and Zinnen, S.
Synthetic ribonucleic acids with RNAse activity
Patent: US 6528640-A 649 04-MAR-2003;
Location/Qualifiers
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AR286277
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            synthetic construct
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Pred. No. 1.6e+02;
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Pred. No. 1.6e+02;
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Patent: WO 0159103-A 2375 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
                                                                                                                              17 bp
Sequence 96 from Patent W00162911.
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Method and reagent for the modulation and dia
nogo gene expression
Patent: WO 0159103-A 2376 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M. (
Location/Qualifiers
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Ellis, J.H. Method and reagent for the inhibition of grid
                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                             Homo sapiens
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/mol type="mRNA"
/db xref="raxon:32630"
/note="Nucleic Acid"
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3 c 8 g
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/mol_type="mRNA"
/db_xref="taxon:32630"
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                                                                                                  Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression,
reversion, apoptosis and/or resistance to viruses and t
                                                                                                                                                                                                                                                      Sequence 1944 from Patent AX673499
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Method and reagent for the inhibition of grid patent: WO 0162911-A 284 30-AUG-2001; RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO GILOCALION/Qualifiers
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                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                          Patent: WO 03004526-A 1944 16-JAN-2003;
                                                                                                                                                                                                          Homo sapiens (human)
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/mol_type="mRNA"
/db_xref="taxon:9606"
6 c 3 g
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/mol_type="mRNA"
/db_xref="taxon:9606"
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/organism="Homo sapiens"
/mol_type="genomic DNA"
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Pred. No. 1.6e+02;
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Pred. No. 1.6e+02;
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Matches 14; Conserv
Query Match
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AX673500
                                                                                                                                                                  Telerman,A., Amson,R. and Tuijnder,M. Sequences involved in phenomena of tumour reversion, apoptosis and/or resistance to
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                                                                                                            Patent: WO 03004526-A 2866 16-JAN-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                                                                      medicines
                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AX674421
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                         /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
4 c 8 g
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/mol_type="genomic DNA"
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Pred. No. 1.6e+02;
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Pred. No. 1.6e+02;
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Pred. No. 1.6e+02;
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AX726400/c
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Matches 14; Conserv
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                     medicines
                                                                                                                                                                                                                                 Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression, tumour version, apoptosis and/or virus resistance and their use
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AX726400
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Location/Qualifiers
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AX727626
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                   Mus musculus (house mouse)
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                                                                                                                    /organism="Mus musculus"
/mol_type="genomic DNA"
/db_xref="taxon:10090"
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/mol_type="genomic DNA"
/db_xref="taxon:10090"
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                                                     Score 13.4; DB 1;
Pred. No. 1.6e+02;
0; Mismatches 1;
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Pred. No. 1.6e+02;
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RESULT 176
AX732843
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AUTHORS
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KEYWORDS
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AX732816/c
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DEFINITION
                                                           Query Match 1.0%;
Best Local Similarity 93.3%;
Matches 14; Conservative
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Best Local
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Sequence 4477 from Patent
AX732843
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Sequence
AX732816
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as medicines.
                                                                                                                                                                                                  Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                                                                                                                                                                medicines
                                                                                                                                                                                                                                                           Telerman,A., Amson,R. and Tuijnder,M. Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as
                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                  Patent: WO 03025175-A 4477 27-MAR-2003;
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Similarity 93.3%;
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4450 from Patent WO03025175.
                                                                                                                      /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
2 c 8 g
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
3 c 6 g
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                                                           Score 13.4; DB 1;
Pred. No. 1.6e+02;
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Pred. No. 1.6e+02;
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RESULT 177 AX739491 LOCUS

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I37420
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I64698
I64698.1
Unclassified.
1 (bases 1 to 17)
Kornbluth, J.
DNA encoding natural killer lytic associated protein
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Draper, K.G., Pavco, P., McSwiggen, J., Gustofson, J. and
Stinchcomb, D.T.
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I37420
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Patent: WO 03025177-A 5081 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
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                                                  Unknown.
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Mammalia; Eutheria;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
3 c 2 g 1
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93.3%;
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Primates;
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Pred. No. 1.6e+02;
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Pred. No. 1.6e+02;
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AR119016
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AR119016
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14; Conserv
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Uchida, K., Uchida, T., Tanaka, Y., Matsuda,
Antisense nucleic acid compound targeted
Patent: US 6150092-A 142 21-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 17)
Draper, K.G., Pavco, P., McSwiggen, J., Gu
Stinchcomb, D.T.
Method of reducing stromelysin RNA via
Patent: US 5731295-A 433 24-MAR-1998;
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Location/Qualifiers
1. .17
                                                                                                                                                               Unknown.
Unclassified.
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RESULT 182

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AR030682/c
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13; Conserv
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1 (bases 1 to 17)
Zwick,M.G., Edington,B.E., McSwiggen,J.A., McGuo,L., Skokut,T.A., Young,S.A., Folkerts,O.
Nucleic acid encoding delta-9 desaturase
                                                                                                        AR196283
Sequence 748 from
AR196283
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Patent: US 5861294-A 18 19-JAN-1999;
Location/Qualifiers
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                                                          Unknown.
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Cowart, M. Daniel.,
                                                                                                                                                                                                                                                                                                                                                                                           Unknown.
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Sequence 18 from patent US
AR030682
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Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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/mol_type="mRNA"
/db_xref="taxon:9606"
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13; Conserv
                                                                               Fattaey,A.R., Jarvis,T., Mcswiggen,J., Booher,R.N. and Holman, Method and reagent for the inhibition of checkpoint kinase-1 (1) enzyme Patent: WO 0157206-A 1033 09-AUG-2001; RIBOZYME PHARMACEUTICALS, INC. (US); Fattaey, Ali R. (US) Location/Qualifiers
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Sequence 1806 from Patent
AX216364
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Location/Qualifiers
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synthetic construct
artificial sequences.
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Patent: WO GISTO 1806 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
MCSwiggen, James (US); Chowrira, Bharat M.
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/note="Nucleic Acid"
/note="C 9 9
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RESULT 188
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Sequence 763 from Patent
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Mammalia; Eutheria; Primates;
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Mammalia; Eutheria; Primates;
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/db_xref="taxon:9606"
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7
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                         767 from Patent EP1281758
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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I37553/c
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                                                1 (bases 1 to 17)
Draper, K.G., Pavco, P., McSwiggen, J.,
Stinchcomb, D.T.
                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 17)
Draper, K.G., Pavco
Stinchcomb, D.T.
Stromelysin targeted ribozymes
Patent: US 5612215-A 567 18-MAR-1997;
Location/Qualifiers
1. .17
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Mammalia; Eutheria; Primates;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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13; Conserv
                                                                                                                                                                                 Method of reducing stromelysin RNA via ribozymes Patent: US 5731295-A 567 24-MAR-1998;
                                                                                                                                                                                                     Unclassified.
1 (bases 1 to 17)
Draper,K.G., Pavco,P., McSwiggen,J., Gustofson,J.
Stinchcomb,D.T.
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Stinchcomb,D.T.
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synthetic construct synthetic construct artificial sequences.
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Sequence 46 from Patent WO0179487.
AX284081
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                                                                  Sequence 47 from Patent AX284082
                                                                                                                                                                                                                                                                                                                                                        Patent: WO 0179487-A 46 25-OCT-2001;
Degitz, Klaus Karl (DE) ; Besch, Robert
Location/Qualifiers
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                                                      AX284082.1 GI:17044792
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Polydesoxyribonucleotides for inhibiting
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Sequenz:Polydesoxyribonukleotid"
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/mol type="genomic DNA"
/db_xref="taxon:32644"
/note="Uncultured Acidobacterium Sub.Div-1"
5 c 6 g 3 t
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                                                                                                                                                                                                                                                                                                     mol_type="genomic DNA"
db_xref="taxon:32630"
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Pred. No. 1.8e+02;
0; Mismatches 2;
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                             Draper,K.G.

Method and reagent for inhibiting he Patent: US 5795778-A 21 18-AUG-1998;

Location/Qualifiers
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AR024071.1
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Sequence 21
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Patent: WO 9104270-A 13 04-APR-1991;
Location/Qualifiers
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synthetic construct
artificial sequences.
1 (bases 1 to 17)
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Degitz, Klaus Karl (DE); Besch, Robert
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
/organism="unknown"
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AR045623/c
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ORGANISM
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AR045623
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Seizinger,B.R., Kley,N.A. and Bianchi,A.B.
NF2 isoforms
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Stinchcomb,D.T. and McSwiggen,J.A.
Interleukin-2 receptor gamma-chain ribozymes
Patent: US 5807743-A 123 15-SEP-1998;
                                                                                                                                                                                                                                                                                                   Sequence 123 from patent AR039275 AR039275.1 GI:5958638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent: US 5872214-A 23 16-FEB-1999;
Location/Qualifiers
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AR036431
Unknown
                  AR045623.1
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1 c 7 g
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+(
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Pred. No. 1.9e+02;
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AR046321
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1 (bases 1 to 17)

Stinchcomb,D.T., Draper,K., McSwiggen,J. and Jarvis,T.

C-myb ribozymes having 2'-5'-linked adenylate residues

Patent: US 5817796-A 416 06-OCT-1998;
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                                                                                                               1 (bases 1 to 17)
Stinchcomb,D.T., Draper,K., McSwiggen,J. and Járvis,T.
Stinchcomb,D.T., Draper,K., McSwiggen,J. and Járvis,T.
C-myb ribozymes having 2'-5'-linked adenylate residues
Patent: US 5817796-A 1114 06-OCT-1998;
Location/Qualifiers
                                                                                                                                                                                     Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unclassified.

1 (bases 1 to 17)

Stinchcomb,D.T., Draper,K., McSwiggen,J. and Jarvis,T.

C-myb ribozymes having 2'-5'-linked adenylate residues

Patent: US 5817796-A 534 06-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unknown
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Stinchcomb.T., Draper,K., McSwiggen,J. and Jarvis,T.
C-myb ribozymes having 2'.5'-linked adenylate residues
Patent: US 5817796-A 2423 06-OCT-1998;
Location/Qualifiers
1 (bases 1 to 17)
Karn, T., Galt, M.John., Heaphy, S. and Dingwall, C. RNA oligonuclectides that bind HIV tat protein Patent: US 5821046-A 18 13-OCT-1998;
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Stinchcomb, D.T., Draper, K., McSwiggen, J. and Jarvis, T.
Stinchcomb, D.T., Draper, K., McSwiggen, J. and Jarvis, T.
Stinchcomb, D.T., Draper, K., McSwiggen, J. and Jarvis, T.
Stinchcomb, D.T., Linked adenylate residues
Patent: US 5817796-A 1116 06-OCT-1998;
Location/Qualifiers
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Karn,J., Gait,M.John., Heaphy,S. and Dingwall,C.RNA oligonucleotides that bind HIV tat protein Patent: US 5821046-A 21 13-OCT-1998;
Location/Qualifiers
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Sequence 19 from patent US
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AR048081
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Karn,J., Gait,M.J., Heaphy,S. and Dingwall,C. Viral (HIV) growth inhibition
Patent: US 6114109_A 18 05-SEP-2000;
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Sequence 18 from patent US
AR108980
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Karn, Gait, M. John., Heaphy, S. and Dingwall, C. RNA oligonucleotides that bind HIV tat protein Patent: US 5821046-A 22 13-OCT-1998;

Location/Qualifiers
                                                              Unclassified.

1 (bases 1 to 17)

Karn, J., Gait, M.J., Heaphy, S. and Dingwall, C. Viral (HIV) growth inhibition

Patent: US 6114109-A 19 05-SEP-2000;
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Karn, J., Gait, M.J., Heaphy, S. and Di
Viral (HIV) growth inhibition
Patent: US 6114109-A 22 05-SEP-2000;
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Karn, Gait, M.J., Heaphy, S. and Di
Viral (HIV) growth inhibition
Patent: US 6114109-A 21 05-SEP-2000;
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                                                                                                   1 (bases 1 to 17)
Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398-A 2761 12-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 17)
Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor recepts
Patent: US 6346398-A 2009 12-FEB-2002;
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Sequence 2761 from patent
AR187273
AR187273.1 GI:20233238
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Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398-A 1992 12-FEB-2002;
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Score 12.8; DB 1;
Pred. No. 1.9e+02;
n. Mismatches 2;
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Pavco,P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 5816 12-FEB-2002;
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1 (bases 1 to 17)

1 (bases 1 to 17)

Paco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J. Method and reagent for the treatment of diseases or crelated to levels of vascular endothelial growth fact Patent: US 6346398-A 4140 12-FEB-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 17)
PavCo, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J. Method and reagent for the treatment of diseases or cond related to levels of vascular endothelial growth factor Patent: US 6346398-A 8106 12-FEB-2002;
                                                                                                                                    Method and reagent for inhibiting herpes simplex virus Patent: US 6440719-A 21 27-AUG-2002;
                                                                                                                                                           1 (bases 1 to 17)
Draper, K.G.
                                                                                                                                                                                                                                               Sequence 21 from patent US AR224290
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Sequence 783 from patent US 6528640.
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El Solh,N. and Allignet,J.
El Solh,N. and their use for detecting resistance to streptogramin A or to streptogramin B and related compounds Patent: US 6506893-A 31 14-JAN-2003;
 Blatt,L.,
                                                                      AX215401.1 GI:15525444
                                                                                 Sequence 843 from Patent
AX215401
                                                                                                                                                                                                                                                                                                         1 (bases 1 to 17)
Beigelman, L. Burgin, A., Beaudry, A., Karpeisky, Matulic-Adamic, J., Sweedler, D. and Zinnen, S. Synthetic ribonucleic acids with RNAse activity Patent: US 5528640-A 783 04-MAR-2003;
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Blatt, L., Mcswiggen, J. and Chowrira, B.M. Method and reagent for the modulation and diagnosis of nogo gene expression Patent: WO 0159103-A 2092 16-AUG-2001; RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, Lawrence McSwiggen, James (US); Chowrira, Bharat M. (US)
                                                                                                                                                                  Sequence 2092 from Patent AX216650
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Sequence 845 from Patent
AX215403
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Patent: WO 0159103-A 843 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US);
MCSwiggen, James (US); Chowrira, Bhaz
Location/Qualifiers
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Patent: WO 0159103-A 845 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M. (
Location/Qualifiers
                                                                                                               synthetic
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/mol_type="mRNA"
/db_xrefs"teaxon:32630"
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AX216661/c
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                                                nogo gene expression
Patent: WO 0519103-A 2341 16-AUG-2001;
RIBDZYME PHARWACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
Location/Qualifiers
                                                                                                              Blatt,L., Mcswiggen,J. and Chowrira,B.M. Method and reagent for the modulation and
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synthetic construct
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McSwiggen, James (US) ; Chowrira,
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/mol type="mRNA"
/db_xref="taxon:32630"
/note="Nucleic Acid"
10 c 5 g 1 t
/organism="synthetic construct"
/mol_type="mRNA";
/db_xref="taxon:32630"
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/mol_type="mRNA"
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AX217829
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AX217484/c
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                                                                                                                 nogo gene expression
Patent: WO 0159103-A 3271 16-AUG-2001;
RIBOZYME PHARWACEUTICALS, INC. (US) ; Blatt,
McSwiggen, James (US) ; Chowrira, Bharat M.
Location/Qualifiers
                                                                                                                                                                                                                                                                                          Sequence 3271 from Patent AX217829
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Patent: WO 0159103-A 2926 16-AUG-2001;
RIBOZYME PHARWACEUTICALS, INC. (US) ; Blatt,
MCSwiggen, James (US) ; Chowrira, Bharat M.
Location/Qualifiers
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                                                                                                                                                                                      Blatt,L., Mcswiggen,J. and Chowrira,B. Method and reagent for the modulation
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Method and reagent for the modulation and diagnosis of cd20 and
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                                                                    organism="synthetic construct"
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Pred. No. 1.9e+02;
         Score 12.8; DB 1;
Pred. No. 1.9e+02;
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AX226704
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                                                                                                                                                                          Patent: WO 0157206-A 76 09-AUG-2001; RIBOZYME PHARMACEUTICALS, INC. (US); Fattaey, Ali R. (US)
                                                                                                                                                                                               Fattaey, A.R., Jarvis, T., Mcswiggen, J., Booher, R.N. and Holman, P.S. Method and reagent for the inhibition of checkpoint kinase-1 (chk lenzyme
                                                                                                                                                                                                                                                                 synthetic construct synthetic construct
                                                                                                                                                                                                                                                                                                                  Sequence 76 from Patent AX226704
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/mol type="mRNA"
/db_xref="taxon:32630"
/note="Nucleic Acid"
7 c 1 g 7 t
                                                                                                /organism="synthetic construct"
/mol type="mRNA"
/db_xref="taxon:32630"
4 c 2 g 5 t
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AX227307/c
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ACCESSION
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AX264368
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AX227715
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Patent: WO 0157206-A 679 09-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US);
LOCATION/Qualifiers
Homo sapiens
Homo sapiens
                                               AX264368
Sequence 1759 from Patent
AX264368
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Sequence 1087 from Patent
AX227715
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                                                                                                                                                                                                                                                                                                           Patent: WO 0157206-A 1087 09-AUG-2001; RIBOZYME PHARMACEUTICALS, INC. (US);
                                                                                                                                                                                                                                                                                                                                                  Fattaey, A.R., Jarvis, T., Mcswiggen, J., Booher, R.N. (Method and reagent for the inhibition of checkpoint
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/mol type="mRNA"
/db_xref="taxon:32630"
4 c 3 g 5 t
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/mol_type="mRNA"
/db_xref="taxon:32630"
.__2 c 0 g 9 t
                                     GI:16513167
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           (human)
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
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Matches 14; Conserv
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Targeted chromosomal genomic alterations stranded oligonucleotides
Patent: WO 0173002-A 1759 04-OCT-2001;
UNIVERSITY OF DELAWARE (US)
Location/Qualifiers
Kmiec, E.B., Gamper, H.B. and Rice, M.C.
Targeted chromosomal genomic alterations with modified single stranded oligonucleotides
Patent: WO 0173002-A 3314 04-OCT-2001;
                                                                                                Homo sapiens (human)
                                                                                                                                                       AX265923
Sequence 3314 from Patent
AX265923
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Mammalia; Eutheria; Primates;
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Mammalia; Eutheria;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
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                                                                    Craniata; Vertebrata; Catarrhini; Hominidae;
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                                                                              Method and reagent for the inhibition of grid Patent: WO 0162911-A 283 30-AUG-2001; RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO G
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Sequence 3315 from Patent
AX265924
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Mammalia; Eutheria; Primates;
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Mammalia; Eutheria; Primates;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
3 c 4 g
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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patent: WO 0188124-A 534 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US) ; GLAXO
Location/Qualifiers
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Sequence 534 from Patent WO0188124.
AX422198
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Sequence 1212 from Patent WO0188124.
AX422876
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="mRNA"
/db_xref="taxon:9606"
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/mol_type="mRNA"
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AX498837
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Mammalia; Eutheria;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="mRNA"
/db_xref="taxon:9606"
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Pred. No. 1.9e+02;
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AX500370/c
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Sequence 1677 from Patent
AX500370
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Human testis expressed
            Zhan, J.
                                    Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                           Homo sapiens
                                                                      Homo sapiens (human)
                                                                                              AX500370.1 GI:23382663
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Mammalia; Eutheria; Primates;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 1.9e+02;
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Sequence 121 from Patent EP1239051.
XX530612
                                                                      Human posh-like protein 1
Patent: EP 1239051-A 124 11-SEP-2002;
Aeomica, Inc. (US)
                                                                                                                                                                                                       Sequence 124 from Patent AX530615
                                                                                                                                 Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
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Patent: EP 1239051-A 121 11-SEP-2002;
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Mammalia; Eutheria;
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                                                                                                           Shannon, M.
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/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Best Local Similarity 87.5%;
Matches 14; Conservative
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                                                                                                                                                                                Human lccl-domain containing protein Patent: EP 1262488-A 701 04-DEC-2002;
                                                                                                                                                                                                                               Homo sapiens (human)
Homo sapiens
Eukaryota, Metazoa, Chordata,
Mammalia, Eutheria, Primates,
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Human lccl-domain containing protein
Patent: EP 1262488-A 700 04-DEC-2002;
Aeomica, Inc. (US)
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Sequence 552 from Patent
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Sequence 589 from Patent
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Mammalia; Eutheria; Primates;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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Sequence 744 from Patent
AX672299
Telerman, A., Amson, R. and Tuijnder, M. Sequences involved in phenomena of to
                                          Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                      Homo sapiens
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Sequence 1108 from Patent AX672663
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent: WO 03004526-A 552 16-JAN-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
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Sequences involved in phenomena of tumour suppression, tumour sersion, apoptosis and/or resistance to viruses and their use as
                                                                                  Homo sapiens (human)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
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Sequences involved in phenomena of tumour suppression,
Sequences involved in phenomena of tumour suppression,
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Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                    Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses and their use
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Sequence 1930 from Patent
AX673485
AX673485.1 GI:29331833
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
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Pred. No. 1.9e+02;
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Sequences involved in phenomena of tumour suppression,
Sequences involved and/or resistance to viruses and I
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                                                                                  Patent: EP 1281758-A 129 05-FEB-2003;
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Location/Qualifiers
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Sequence 3229 from Patent
AX674784
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Similarity 87.5%;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/organism="Homo sapiens"
/mol_type="genomic DNA"
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
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AX687586
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 130 from Patent EP1281758.
AX687398
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Four human zinc-finger-containing proteins
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Homo sapiens
Eukaryota, Metazoa, Chordata,
Mammalia, Eutheria, Primates,
                                                                                                                                                                                               Patent: EP 1281758-A 318 05-FEB-2003;
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Similarity 87.5%;
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GGCTCCAGGAGCTCTG 17
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Best Local Similarity 87.5%;
Matches 14; Conservative
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Best Local Similarity
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Patent: EP 1281758-A 341 05-FEB-2003;
                                                                                                                                                                                                                                                                                                Shannon, M., Gu, Y. and Nguyen, C.T. Four human zinc-finger-containing proteins
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 341 from Patent EP1281758. AX687609
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Sequence 320 from Patent EP1281758.
AX687588
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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                                    Homo sapiens (human)
                                                                                          Sequence 1130 from Patent AX688398
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            Mammalia;
                                                                             AX688398.1 GI:29411100
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                         Eukaryota;
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Mammalia; Eutheria;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                     Sequence 5250 from Patent AX692518
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              Aeomica,
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                                                               Shannon, M.,
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                                                  Four human zinc-finger-containing proteins
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AX722358
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ACCESSION
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                                                                                                                                                         Telerman,A., Amson,R. and Tuijnder,M. Sequences involved in phenomena of tumour suppression,
                                                                                                                                                                                                                                                                              17 bp
Sequence 45 from Patent WO03025176.
AX722358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Method and reagent for inhibiting HBV viral replication Patent: EP 1288296-A 355 05-MAR-2003; RIBOZYME PHARMACEUTICALS, INC. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Herpes simplex virus unknown type
Herpes simplex virus unknown type
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 355 from Patent AX711055
                                                                                           Molecular Engines Laboratories
Location/Qualifiers
                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                                                   AX722358.1 GI:30422859
                                                                                                                     Patent: WO 03025176-A 45 27-MAR-2003;
                                                                                                                                    medicines
                                                                                                                                              reversion, apoptosis and/or virus resistance
                                                                                                                                                                                                                                         Mus musculus (house mouse)
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                                       /organism="Mus musculus"
/mol_type="genomic DNA"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Herpes simplex virus
/mol_type="genomic RNA"
/db_xref="taxon:126283"
5 c 7 g
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/db_xref="taxon:9606"
1 c 3 g
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Pred. No. 1.9e-
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  Score 12.8;
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Pred. No. 1.9e+02;
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AX723196/c
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ACCESSION
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AX722882
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1849 AAAGACCTTTCTGATC 1864
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14; Conserv
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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AX722882
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                                                                                                                                                       Patent: WO 03025176-A 883 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                                                                                                                      Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression, tumour suppression, apoptosis and/or virus resistance and their use
                                                                                                                                                                                                                                                                                                                                         Sequence 883 from Patent AX723196
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Sequences involved in phenomena of tumour suppression, reversion, apoptosis and/or virus resistance and their
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:10090"
2 c 6 g
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                                                                                                                organism="Mus musculus"
/mol_type="genomic DNA"
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                                     Score 12.8; DB 1;
Pred. No. 1.9e+02;
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0; Mismatches 2;
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Pred. No. 1.9e+02;
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WO03025176
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WO03025176.
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Sciurognathi; Muridae;
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AX724673/c
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AX724155/c
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RESULT 277 AX724703
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TITLE
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                                                                                                                              Query Match
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                                                                                                                                                                                                                                          Patent: WO 03025176-A 2360 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as
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Sequences involved in phenomena of tumour suppression, tumoreversion, apoptosis and/or virus resistance and their use
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Sequence 1842 from Patent
AX724155
                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus
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Similarity 87.5%;
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Mammalia; Eutheria; Rodentia;
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Similarity 87.5%;
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                                                                                                                                                                       /organism="Mus musculus"
/mol_type="genomic DNA"
/db_xref="taxon:10090"
6 C 4 g
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/db_xref="taxon:10090"
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Pred. No. 1.9e+02;
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AX725084
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           Sequence 3464 from Patent AX725777 AX725777.1 GI:30505120
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Location/Qualifiers
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Mus musculus
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                     AX725777
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/mol_type="genomic DNA"
/db_xref="taxon:10090"
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/mol_type="genomic DNA"
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AX726992
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Sequences involved in phenomena of tumour suppression, tumo
Sequences involved in phenomena of tumour suppression, tumo
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                                     Mus musculus
           Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                Mus musculus (house mouse)
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Sequence 3556 from Patent
AX725869
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Location/Qualifiers
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/mol_type="genomic DNA"
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        Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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AX727138
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Best Local Similarity
Matches 14; Conserv
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Best Local Similarity 87.5%;
Matches 14; Conservative
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TITLE
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Sequence
AX727199
Telerman,A., Amson,R. and Tuijnder,M. Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as medicines
Patent: WO 03025176-A 4886 27-MAR-2003;
                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Telerman, A., Amson, R. and Tuljnder, M.
Sequences involved in phenomena of tumour suppression,
Sequences involved and/or virus resistance and their
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Sequence 4825 from Patent WO03025176.
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                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                   Mus musculus (house mouse)
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Location/Qualifiers
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
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Mus musculus
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Pred. No. 1.9e+02;
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AX728341/c
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AX727780
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Matches
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Best Local Similarity
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                                                                                               Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression, reversion, apoptosis and/or virus resistance and their
                                                                                                                                                                                                                                               Sequence 6028 from Patent AX728341
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Location/Qualifiers
                                         Molecular Engines Laboratories
Location/Qualifiers
                                                                                                                                                          Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Mammalia; Eutheria; Rodentia;
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                                                                       Patent: WO 03025176-A 6028 27-MAR-2003;
                                                                                    medicines
                                                                                                                                                                                                      Mus musculus (house mouse)
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/db_xref="taxon:10090"
3 c 2 g
/mol_type="genomic DNA"
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/db_xref="taxon:10090"
3 c 6 g
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                organism="Mus musculus"
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Pred. No. 1.9e+02;
0; Mismatches 2
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Pred. No. 1.9e+02;
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AX728590/c
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Query Match
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Sequence
AX728422
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Sequence 224 from Patent
AX728590
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Sequences involved in phenomena of tumour suppression, tumo
reversion, apoptosis and/or virus resistance and their use
                                                                                                                       Molecular Engines Laboratories (FR)
                                                                                                                                                                reversion, apoptosis and/or virus resistance
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                 Homo sapiens
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                         /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Sequences involved in phenomena of tumour suppression,
Sequencian, apoptosis and/or virus resistance and their
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Molecular Engines Laboratories (FR)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
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Sequence 930 from Patent WO03025175.
AX729296
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Sequence 960 from Patent
AX729326
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Sequences involved in phenomena of tumour suppression, tumo reversion, apoptosis and/or virus resistance and their use
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 2026 from Patent
AX730392
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Sequences involved in phenomena of tumour suppression, tumour suppression, apoptosis and/or virus resistance and their use
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Sequence 1214 from Patent
AX729580
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AX729354
AX729354.1 GI:30508697
Homo sapiens (human)
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Mammalia; Eutheria; Primates;
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Mammalia; Eutheria; Primateß; Catarrhini; Hominidae; Homo.
                             AX730392.1 GI:30509735
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Molecular Engines Laboratories (FR)
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                         Homo sapiens (human)
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Location/Qualifiers
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 3121 from Patent
AX731487
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Telerman,A.,
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/mol_type="genomic DNA"
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Amson, R. and Tuijnder, M.
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Location/Qualifiers
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Patent: WO 03025175-A 3682 27-MAR-2003;
Molecular Engines Laboratories (FR)
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                                          Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suj
                                                                                                     Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                   Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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Local Similarity 87.5%;
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                                                                                                               Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as
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Sequence 5107 from Patent
AX733473
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                                                         Molecular Engines Laboratories (FR)
Location/Qualifiers
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Molecular Engines Laboratories (FR)
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Sequence 3683 from Patent
AX732049
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 1.9e+02;
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AX733774/c
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Sequences involved in phenomena of tumour suppression, tumour suppression, apoptosis and/or virus resistance and their use
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Sequence 5584 from Patent
AX733950
                                                                                                                                      Patent: WO 03025175-A 5584 27-MAR-2003;
Molecular Engines Laboratories (FR)
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Molecular Engines Laboratories (FR)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                     /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Eutheria; Primates;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 357 from Patent
AX734767
                                                                                                                                                                          Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression, reversion, apoptosis and/or resistance to viruses and thereof as medicaments
Patent: WO 03025177-A 468 27-MAR-2003;
Molecular Engines Laborateories (FR)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                      Sequence 468 from Patent AX734878
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Patent: WO 0302517-A 357 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
5 c 3 g
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/mol_type="genomic DNA"
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AX738552 17 bp DNA Sequence 4142 from Patent W003025177.
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Patent: WO 03025177-A 3051 27-WAR-2003;
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses and the use thereof as medicaments patent: WO 03025177-A 768 27-MAR-2003; Molecular Engines Laboratories (FR)
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Similarity 87.5%;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
3 c 5 g
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
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AX738641/c
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Sequence 492 from Patent
AX744527
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Location/Qualifiers
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Patent: WO 03025177-A 4142 27-MAR-2003;
Molecular Engines Laboratories (FR)
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Sequences involved in phenomena of tumour suppression, tun
reversion, apoptosis and/or resistance to viruses and the
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AX738552.1 GI:30517840
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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                GI:30723194
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ORGANISM

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Homo sapiens (human) Homo sapiens

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                                                                                                                                                                                                                                                                                                                                                                                                                                         A human G protein coupled receptor
Patent: WO 03031621-A 494 17-APR-2003;
Amersham Biosciences (GV) Corp. (US)
Location/Qualifiers
RIBOZYME PHARMACEUTICALS INC OS Artificial Sequence PN JP 2000342285-A/335
                                     Draper, K.G., Dadyktz, L.W., Macswigen, J.A., Maysejak, D.G., Holsesk, J.J. and Mamone, A.J. Matched and reagent for inhibiting viral replication Patent: JP 2000342285-A 335 12-DEC-2000;
                                                                                                                                             BD001175.1 GI:18625734
JP 2000342285-A/335.
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AX744529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AX744529.1 GI:30723196
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/db_xref="taxon:9606"
8 c 1 g
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7 c 1 g
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                TITLE
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Best Local Similarity 87.5%;
Matches 14; Conservative
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PF 01-MAY-1992 US 07/882689,14-MAY-1992 US

14-MAY-1992 US 07/882713,14-MAY-1992 US

14-MAY-1992 US 07/88283,14-MAY-1992 US

14-MAY-1992 US 07/882886,14-MAY-1992 US

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14-MAY-1992 US 07/883849,14-MAY-1992 US

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14-MAY-1992 US 07/884492,14-MAY-1992 US

14-MAY-1992 US 07/884436,14-MAY-1992 US

15-OCT-1992 US 07/983322,14-MAY-1992 US

15-OCT-1992 US 07/963322,07-DEC-1992 US

17-DEC-1992 US 07/963322,07-DEC-1992 US

17-DEC-1992 US 07/963322,07-DEC-1992 US

18-CENNETH G DRAPER, LEC W DADYKTZ, JAMES A MACSWI
                                                                                                                                                                        RIBOZYME PHARMACEUTICALS I
OS Artificial Sequence
PN JP 2000342286-A/335
PD 12-DEC-2000
PF 01-MAY-2000 JP 200013
                                               14-MAY-1992
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PI JAMES
PC C12N15/00,
C12N15/00,
PC C12N5/
CC C12N5/
FH Key
FT BOUXCE
                                                                                                                                                                                                                                               synthetic construct
synthetic construct
artificial sequences.

1 (bases 1 to 17)
1 (bases 1 to 17)
Draper, K.G., Dadyktz, L.W., Macswigen, J.A., Maysejak, D.
Holseek, J.J. and Mamone, A.J.
Method and reagent for inhibiting viral replication
Patent: JP 200342286-A 335 12-DEC-2000;
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JP 2000342286-A/335
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1Y-1992 US
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C12N15/09, C12N5/10, C12N7/00, C12N9/22//(C12N5/10, C12R1:91),
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/mol_type="genomic RNA"
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100 JP 2000132651

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101 JP 2000132689,14-MAY-1992 US

S 07/882713,14-MAY-1992 US

S 07/882882,14-MAY-1992 US

JS 07/882889,14-MAY-1992 US

US 07/882899,14-MAY-1992 US

US 07/882922,14-MAY-1992 US

US 07/883849,14-MAY-1992 US

US 07/88384074,14-MAY-1992 US

US 07/8844074,14-MAY-1992 US

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BD067344/c
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KEYWORDS
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15-OCT-1992 US 07/9
07-DEC-1992 US 07/9
KENNETH G DRAPER, LEC W
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PI JAME
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PC A61F
PC A61P
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Akhtar,S., Fell,P. and Mcswiggen,J.A.
Akhtar,S., rell,P. and Mcswiggen,J.A.
Enzymatic nucleic acid treatment of diseases or conditions related to levels of epidermal growth factor receptors
Patent: JP 2001511003-A 184 07-AUG-2001;
                                                                                                           3338
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                                                                                                                                                                                                                                                                                                                                                                                                            unidentified
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source
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Enzymatic nucleic acid treatment related to
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                                                                                                                                                                                                                                           14-JAN-1998 JP 1998532913
31-JAN-1997 US 60/0364
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C12N15/09,C12N5/10,C12N7/00//A61K38/43,A61K39/125,A61K39/13,
A61K39/135,
A61K39/145,A61K39/21,A61K39/23,A61K39/245,A61K39/29,A61K48/00,
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2N15/09,C12R1:93),C12N15/00,C12N5/00,A61K37/48,(C12N15/00, )
                                                                                                                                                                                                  Strandedness: Single;
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                                                                                                                               levels of epidermal growth factor receptors
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                                                                                           /organism='Unidentified'
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/db_xref="taxon:32630"
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07/93322,07-DEC-1992 US 07/9871
07/987130,07-DEC-1992 US 07/9871
JEC W DADYKTZ,JAMES A MACSWIGEN, PI
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BD104906/c
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PR 31-JAN-1997 US 60/036476,04-DEC-1997 US SAGHIR AKHTAR, PATRICIA FELL, JAMES A MCSWIGGEN PC C12N9/00,C07K14/71
CC Strandedness Single;
CC Topology: Linear
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Kit and method for determining HLA type
Patent: WO 0192572-A 1010 06-DEC-2001;
NISSHINBO INDUSTRIES INC, SYSTEM RESEARCH INC, HIDETOSHI INOKO,
                                      Inoko,H., Kagiya,T., Ichihara,T., Matsumura,Y.,
Nishida,M.
                                                                                                    BD104906.1 GI:22650480
WO 0192572-A/1010.
synthetic construct
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Akhtar,S., Fell.P. and Mcswiggen,J.A.
Enzymatic nucleic acid treatment of diseases or conditions related
to levels of epidermal growth factor receptors
Patent: JP 2001511003-A 326 07-AUG-2001;
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JP 200151100
                                                               artificial sequences.

1 (bases 1 to 17)
                                                                                         synthetic construct
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Enzymatic nucleic acid treatment of diseases
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Location/Qualifiers
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/mol_type="genomic RNA"
/db_xref="taxon:32644"
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TATSUO ICHIHARA, YOSHIYUKI MATSUMURA, SHOGO

MORIYA, MICHIO

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PI SHOGG
PC C12Q1
CC Desci
FH Key
FT sourc
                                                                                                                                                                                                                                                                                                              1 (bases 1 to 17)
Fuchs, R.L., Kishore, G.M. and MacIntosh, S.C.
Method for improving the efficacy of insect
Patent: EP 0339009-A2 9 25-OCT-1989;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PR PR PR
                              1 (bases 1 to 17)
Seizinger, B.R., Kley, N.A. and Bianchi, A.B.
                                                                                                                                                                                                                                                                                                                                                                                  Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NISHIDA
            Patent: US 5578462-A 23 26-NOV-1996;
                                                      Unclassified.
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W0 0192572-A/1010
06-DEC-2001
01-JUN-2001 W0 2001JP004662
01-JUN-2000 JP 00P 164798
HIDETOSHI INOKO,TAEKO KAGIYA,TATSUO ICHIHARA,YOSHIYUKI
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C12Q1/68, C12M1/00, C12N15/09, G01N33/53
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Location/Qualifiers
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7 c 2 g
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I37681
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Draper.K.G., Pavco,P., McSwiggen,J., Stinchcomb,D.T.
Stromelysin targeted ribozymes
Patent: US 5612215-A 695 18-MAR-1997;
                                                                                                                                                                                Unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Unclassified.
1 (bases 1 to 17)
Draper, K.G., Pavco, P., McSwiggen, J., Stinchcomb, D.T.
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152675
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152793
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Stinchcomb, D.T., Draper, K., McSwiggen, J.
C-myb targeted ribozymes
Patent: US 5646042-A 534 08-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 17)
Stinchcomb, D.T., Draper, K., McSwiggen, J.
C-myb targeted ribozymes
Patent: US 5646042-A 416 08-JUL-1997;
Location/Qualifiers
                                       1 (bases 1 to 17)
Stinchcomb.D.T., Draper,K., McSwiggen,J.
C-myb targeted ribozymes
Patent: US 5646042-A 1114 08-JUL-1997;
                                                                                                                                      Sequence 1114 from patent I53373
                                                                                                                                                                                                                                                                                                                                                                                                        Unknown.
                                                                                    Unclassified.
                                                                                               Unknown
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                                                                                                                            153373.1
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IS3375
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Stinchcomb, D.T., Draper, K., McSwiggen, J.
C-myb targeted ribozymes
Patent: US 5646042-A 2423 08-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 17)
Stinchcomb, D.T., Draper, K., McSwiggen, J.
C-myb targeted ribozymes
Patent: US 5646042-A 1116 08-JUL-1997;
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Pred. No. 1.9e+02;
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Draper, K.G., Pavco, P., McSwiggen, J., Gustofson, J
Stinchcomb, D.T.
Method of reducing stromelysin RNA via ribozymes
Patent: US 5731295-A 695 24-MAR-1998;
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Draper, K.G., Pavco, P., McSwiggen, J., Gustofson, J.
Stinchcomb, D.T.
Method of reducing stromelysin RNA via ribozymes
Patent: US 5731295-A 694 24-MAR-1998;
                                                                                     of fowlpox viruses
Patent: US 6103244-A 75 15-AUG-2000;
                                                                                                                                                                                                     Sequence 75 from patent US
AR105805
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Dorner, F., Scheiflinger, F.,
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1 (bases 1 to 14)

Sykes, K.F. and Johnston, S.A.

Methods of screening open reading frames to determine whether they encode polypeptides with an ability to generate an immune response Patent: US 6410241-A 3 25-JUN-2002;
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PATENE: JP 2002017374-A 5 22-JAN-2002;
MINEO ISEKI MASAKATSU WATANABE
OS Artificial Sequence
PN JP 2002017374-A/5
PD 22-JAN-2002
PF 03-JUL-2000 JP 2000240426
PI MINEO ISEKI MASAKATSU WATANABE
PC C12N15/09.C12N9/88//(C12N9/88,C12R1:89)
unidentified unidentified unclassified.
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A40008 GI:2296192
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Sequence 13 from Patent WO9526410.
A46520
Other publication AU 1444397 19970820
            Stuyver,L., Louwagie,J. and Rossau,R. METHOD FOR DETECTION OF DRUG-INDUCED MUTATIONS TRANSCRIPTASE GENE Patent: WO 9727332-A 157 31-JUL-1997; INNOGENETICS NV (BE)
                                                                                                                                                           A64369
Sequence 157 from Patent
A64369
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Windass, J. D.
BIOLOGICAL INSECT CONTROL AGENT
Patent: WO 9526410-A 13 05-OCT-1995;
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Guest, P.J., Windass, J.D., Suner, M. and Earley, F.G.
BIOLOGICAL CONTROL AGENTS CONTAINING MOLLUSC TOXINS
Patent: WO 9423047-A 13 13-OCT-1994,
ZENECA LTD (GB)
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13; Conserv
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Schreiber, A.D. and Park, J.-G.
Method of inhibiting phagocytosis
Patent: US 5858981-A 31 12-JAN-1999;
Location/Qualifiers
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AR028992
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Schreiber, A.D. and Park, J.-G.
Method of inhibiting phagocytosis
Patent: US 5858981-A 27 12-JAN-1999;
                                                                                                                                                                      Unclassified.
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
5 c 1 g
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
                                        Score 12.4; DB 1;
Pred. No. 1.9e+02;
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RESULT 337

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AR156870/c
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AR133690/c
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Sequence
AR102668
                             1 (bases 1 to 15)
Schreiber, A.D. and Park, J.-G.
Methods of inhibiting phagocytosis
Patent: US 6242427-A 27 05-JUN-2001;
                                                                                     Unknown
                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 15)
Stinchcomb,D.T., Jarvis,T. and McSwiggen,J.
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 2115 27-FEB-2001;
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Patent: US 6087093-A 157 11-JUL-2000;
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Method for detection of drug-induced mutations
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/organism="unknown"
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AR262971/c
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SOURCE
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AR192985
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                                                                                                                                                                                                1 (bases 1 to 15)
PavCo,P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 8473 12-FEB-2002;
                                                                                                                                                                                                                                                                                                        AR192985
Sequence 8473 from patent
AR192985
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 15)
Schreiber, A.D. and Park, J.-G.
Methods of inhibiting phagocytosis
Patent: US 6242427-A 31 05-JUN-2001;
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                                                                                                                                                                                                                                                                Unknown.
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THIRD WAVE TECHNOLOGIES, INC. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13;
Brower, A., Brow, M.A., Cracauer, R.F., Fors, L., Granske, R., de arruda Indig, M., Kurensky, D., Luedtke, C., Lukowiak, A.A., Lyamichev, V., Neri, B.P., Reimer, N.D., Roeven, R.T., Skrzypczynski, Z., Ziarno, W.A., Comerford, J., Stump, S. and Viegut, D.D.
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                                                                                                                                       AX598514
Sequence 788 from Patent
AX598514
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artificial sequences.
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Sequence 786 from Patent
AX598512
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                                                                                                                           AX598514.1
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5 c 1 g
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/db_xref="taxon:32630"
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                                                                                                                           GI:28398692
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Pred. No. 1.9e+02;
0; Mismatches 1
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Pred. No. 1.9e+02;
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AX637907
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                                   1 (bases 1 to 15)
Yabusaki,Y., Murakami,H., Sakaki,T., Shibata,M.
SOLUBLE NADPH-CYTOCHROME P-450 REDUCTASE
Patent: JP 1994113836-A 5 26-APR-1994;
AGENCY OF IND SCIENCE & TECHNOL
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Patent: EP 1260586-A 5046 27-NOV-2002;
RIBOZYME PHARMACEUTICALS, INC. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Systems and method for detection assay production and sale patent: WO 0244994-A 788 06-JUN-2002; THIRD WAVE TECHNOLOGIES, INC. (US)

Location/Qualifiers
8 0 S
                                                                                                                                                                            Synthetic DNA linkers
E07289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stinchcomb,D.T., Dudycz,L.W., Chowrira,B., Grimm,S., Direnzo
Karpeisky,A., Draper,K.G., Kisich,K., Matulic-Adamic,J.,
Mcswiggen,J.A., Modak,A., Pavco,P., Beigelman,L., Sullivan,S
Sweedler,D., Thompson,J.D., Tracz,D., Usman,N., Wincott,F.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 5046 from Patent AX637907
                                                                                                                                                JP 1994113836-A/5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   unidentified unidentified
                                                                                                       unclassified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Method and reagent for inhibiting the expression of disease related
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            Artificial sequences.
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/db_xref="taxon:32644"
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[mol_type="genomic DNA"]
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Pred. No. 1.9e+02;
0; Mismatches 1
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Pred. No. 1.9e+02;
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Sullivan,S.M.,
                                                                               Okawa, H.
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                                                                                                                    OS None
OC Artificial
PN JP 1996228
PD 10-SEP-199
PP 12-AUG-198
PF YABUSAKI YV
SHIBATA MEGUMI,
PI OKAWA HIDEK
PC C12N15/09/
PC C12R11865)
PC C12R11865)
PC C12R11865)
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PC T2R11865)
PC Strandednes
CC topology:
PT SOURCE
FH Key
FT SOURCE
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PI OKAWA HIDE
PC C12N9/02,C
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Yabusaki, Y., Murakami, H., Sakaki, T., Shibata, M.
PRODUCTION OF CHIMBRA OXIDASE
Patent: JP 1996228776-A 4 10-SEP-1996;
AGENCY OF IND SCIENCE & TECHNOL
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25-OCT-1991 JP 1991305592
YABUSAKI YOSHIYASU, MURAKAMI HIROKO,
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topology: Linear;
hypothetical: No;
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hypothetical: No;
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C12N15/09//C12N1/19,C12N9/02,(C12N1/19,C12R1:865),(C12N9/02
                                                                                                                                                                                                                                                                 YABUSAKI YOSHIYASU,
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10-SEP-1996
12-AUG-1986 JP 1995347122
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                                     /organism="unidentified"
/mol_type="genomic_DNA"
/db_xref="taxon:32644"
5 c 3 g
                                                                                                     /organism='Artificial sequences'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mol_type="genomic DNA"
/db_xref="taxon:32644"
5 c 3 g
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Location/Qualifiers
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0.9%;
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Score 12.4; DB 1;
Pred. No. 1.9e+02;
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Pred. No. 1.9e+02
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            Length
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AX132931/c
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AR054087/c
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         Homo sapiens (human)
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Goldenberg,T and Tritz,R.
Ribozyme therapy for the inhibition Patent: US 5834440-A 14 10-NOV-1998;
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                                               Sequence 4149 from Patent
AX132931
                                                                                                                                                                                                                                                                                                                       Sequence 14 from AR054087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 15)
McSwiggen, J., Draper, K., Pavco, P. and
Respiratory syncytial virus ribozymes
Patent: US 5693532-A 296 02-DEC-1997;
Eukaryota; Metazoa; Chordata; Craniata;
                                       AX132931.1
                                                                   AX132931
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Vertebrata; Euteleostomi;
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                                                                         Homo sapiens (human)
                                                                                                                             Sequence 29 from Patent AX320908
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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                         Tian, H.,
                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                               AX320908.1 GI:17902457
                                                                                                                                                      AX320908
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                         Schultz, J. and Shan, B.
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/db_xref="taxon:9606"
/note="Hairpin ribozyme recognition site for PCNA"
4 c 1 g 7 t
                                                                                                                                                                                                                                                                                                                          /note="Hammerhead ribozyme recognition site
4 c 1 g 7 t
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Catarrhini;
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Search completed: December 1, 2003, 11:51:24 Job time : 6 secs
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Sequence 25 from Patent W002095071.
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                                                                                                                                                                                                                                                                                   Means and methods for identifying genes and proteins involved in the prevention and/or repair of a replication error Patent: WO 02095071-A 25 28-NOV-2002;
Koninklijke Nederlandse Akademie van Wetenschappen (NL)
Location/Qualifiers
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synthetic construct
artificial sequences.
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/mol type="genomic DNA"
/mol type="genomic DNA"
/db_xref="taxon:9606"
/note="5' splicing site for exon 6"
/note="5' splicing site for exon 6"
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Kinase domain 3' P

Mouse fibroblast g

FGF receptor gene

Downstream primer

PCR primer used to

FGFR-2 (bek) PCR

A KIT gene primer KI

KIT gene PCR forwa

VEGFR-1 gene specif

Human VEGFR-1 RT-P

Rat FGFR coding se

Primer #2 for huma

PCR primer #3 to i

Primer for SCP an

Human DP2.5 APC pr

Primer for SCP an

Human APC (DP2.5)

DP1, SRP19, DP25

Primer #2 for huma

PTK 6 gene specifi

PTK 10 gene specifi

PTK 14 gene specifi

PTK 16 gene specifi

PTK 17 gene specifi

PTK 18 gene specifi

PTK 19 gene specifi
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Kinase domain 3' P
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ID AAQ13309;
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AAQ13309;
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AAQ13309;
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25-MAR-2003 (updated)
DT 28-OCT-1991 (first entry)
XX
Probe OAB984 for bFGF rece
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Basic fibroblast growth fa
XX
OS Synthetic.
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WO9111459-A.
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C1-JAN-1991; 91WO-EP0010
XX
PD 08-AUG-1991.
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PD 08-AUG-1991; 90GB-000144
XX
PR 23-JAN-1990; 90GB-00014
PR 23-JAN-1990; 90GB-00144
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PR 23-JAN-1991; 91WO-EP0010
XX
WFI; 1991-252611/34.
XX
PR WPI; 1991-252611/34.
XX
PR WPI; 1991-252611/34.
XX
PR Extracellular form of huma
PT used to treat tumours, abb
PT retinopathy, rheumatoid as
PT contraceptives.
XX
Example 1; Page 11; 29pp;
XX
The probe was used to scre
CC library for the gene encoor
CC from the partial CDNA Clor
CC See also AAQ13308-Q13311.
CC (Updated on 25-MAR-2003 to
XX
SQ Sequence 30 BP; 5 A; 8 C;
RESULT 2
AAF58623
ID AAF5
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AAF AAF5
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AC AAF5
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Matches 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Extracellular form of human fibroblast growth factor receptor used to treat tumours, abnormal angiogenesis e.g. diabetic retinopathy, rheumatoid arthritis and arteriosclerosis and as contraceptives.
                            WO200107627-A1
                                                                              Mouse; c-kit; drosophila recombination associated protein; DRAP; gene targeting; ss.
                                                                                                                         Murine c-kit exon 17 potential mutagenic oligonucleotide
                                                                                                                                                   27-APR-2001
                                                                                                                                                                              AAF58623;
                                                                                                                                                                                                       AAF58623 standard; cDNA; 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 11; 29pp; English
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93.1%;
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Length 30;

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RESULT 3
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ID AAV440
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                        Assays for high-affinity heparin-binding growth factor receptor ligands - using receptor-overexpressing cells or cell-free syst
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Basic fibroblast growth factor receptor; bFGF; heparin binding; murine; antitumour agent; inhibitor; wound healing; PCR primer; ss.
                                                                                                         WPI; 1998-446074/38.
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14-DEC-1993;
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01-OCT-1998
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                                                                                                                                                          Flanagan JG,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The DRAP gene is suitable for very efficient gene targeting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (YESH ) UNIV YESHIVA EINSTEIN COLLEGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2137
                                                                                                                                                                                                               CHILDRENS MEDICAL CENT HARVARD COLLEGE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8; Page 7; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TGTATTCATCGAGATTTAGCAGCCAGAAAT 2166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGTATTCACAGAGATTTGGCAGCCAGGAAT 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (updated)
(first entry)
                                                                                                                                                        Klagsbrun M,
                                                                                                                                                                                                                                                                                            90US-0631717.
93US-0166717.
                                                                                                                                                                                                                                                                                                                                                                            93US-0166717.
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86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                             Leder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 23.6;
Pred. No. 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                Ornitz
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K
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 31;
                                                                                                                                                          Yayon
                        system
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AAV05314/c
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Best Local
         This oligonucleotide was designed to amplify the region of homology in the kinase domains of PDGF receptor, HER2, HER3, FGF-FLG, FGF-BEK, insulin receptor and IRR. It was used with another kinase homology domain-based primer (see AAV05313) in the amplification of human LIM-kinase 1 (LIMK1) sequences. The LIMK1 gene is composed of 16 exons (see AAV05315 and AAT99599-T99629) and located 15.4 kb 3 of elastin in chromosome 7. It encodes a novel protein kinase (see AAW46576). Williams syndrome cognitive profile (WSCP) is detected by determining zygosity of the LIMK1 locus, with hemizygosity being indicative of impaired visuo-spatial constructive cognition. Chromosome 7 deletion analysis allows
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV44043-V44045 are PCR primers used to amplify a murine basic fibroblast growth factor (bFQF) which is a member of the heparin-binding growth factor receptor family. This protein is used in a method which assays the ability of a substance to bind to a high-affinity heparin-binding growth factor (HBGF) receptor. The assay screens for potential antitumour agents that inhibit binding of HBGF to high-affinity receptors, or for potential wound healing agents that promote such binding.
                                                                                                                                                                                                                                              Diagnosing Williams syndrome cognitive profile from hemizygosity of LIMK1 - gene on chromosome 7 encoding new kinase, allowing differentiation from classic Williams syndrome and supra-vascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Williams syndrome cognitive profile; WSCP; cognition; LIM-kinase 1; LIMK1 gene; supra-vascular aortic stenosis; protein kinase; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV05314;
                                                                                                                                                                                                 Example 3; Page 22; 62pp; English
                                                                                                                                                                                                                                 aortic
                                                                                                                                                                                                                                                                                                             WPI; 1998-101185/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                      07-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV05314 standard;
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                                                                                                                                                                                                                                                                                                                                          Keating MT,
                                                                                                                                                                                                                                                                                                                                                                          (UTAH ) UNIV UTAH RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                       10-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-JAN-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9801740-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kinase domain 3' PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (Updated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GAGATGGAGATGAAGATGAAT 5
                                                                                                                                                                                                                                                                                                                                          Morris CA;
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between WSCP,
                                                                                                                                                                                                                                                                                                                                                                                                                                 97MO-US11687.
                                                                                                                                                                                                                                                                                                                                                                                                       96US-0678039
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.6%;
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Pred. No. 1
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RESULT 6
AAT63286/c
ID AAT63286 standard; DNA; 21
XX
AC AAT63286;
XX
DT 21-MAY-1997 (first entry)
XX
DE FGF receptor gene downstre
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 5
AAQ527728/c
ID AAQ5277
XX AAQ527
XX Pibrob
XX Pibrob
XX Synthe
XX Synthe
XX US5270
XX OS5270
XX OS5270
XX HAPD
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PT Seltes
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Best Local S
Matches 22
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Best Local Similarity
Matches 22; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 27 BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          factor cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   conserved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This primer and its 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Column 7; 37pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                agents or wound healing promoters
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cells having high-affinity heparin-binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1993-404932/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Klagsbrun M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CHIL-) CHILDRENS MEDICAL CENT (HARD ) HARVARD COLLEGE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-DEC-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-DEC-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-DEC-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US5270197-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fibroblast growth factor; DNA primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse fibroblast growth factor 3' DNA primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ52728;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ52728 standard; DNA; 27
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                                                                                                                                                                                                                                                                                                                                              1874
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2269 CCAGTCAAGTGGATGGCTCCAGA 2291
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            are used for screening substances
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                                                                                                                                                                                                                                                                                                                               AGATGGAGATGATGAAGATGATT 1896
                                                                                                                                                                                                                                                                                                    AGATGGAGATGATGAAGATGAAT 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           er and its 5' partner (AAQ52727) correspond to regions highly among mouse BEK, human FLG and chicken fibroblast growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 4 A; 10 C;
                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Leder P,
                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                           downstream primer binds bases 110-130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 10 C; 1 G; 10 T; 0 other;
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95.7%;
                                                                                                                                                                                                                                                                                                                                                                                                             1.5%;
                                                                                                                                                                         21 BP
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Pred. No. 18;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                         Score 21.4; D
Pred. No. 21;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       g growth factor binding for e.g. anti-tumour
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
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The invention relates to methods for promoting corneal cell proliferation in vivo by treating the cells with hepatocyte growth factor (HGF) and optionally keratinocyte growth factor (KGF). Methods for suppressing corneal cell growth include administering Ca ions to the cells. The methods are used for the treatment of corneal disorders following accidental injury, ocular surgery or due to corneal disorders caused by abnormal healing processes of the corneal epithelium and endothelium. The methods are based on the discovery that corneal tissue can express mRNA for HGF, KGF and their respective receptors. The primers AAT63283 and AAT83286 were used to amplify a 205 fragment of the fibroblast growth factor (FGF) receptor 2 cDNA. This primer is the fibroblast growth factor (FGF) receptor 2 cDNA. This primer is the pareful of the primers primer is the methods are based on the second to bases 110-130 of the pareful of the methods of the primers and corresponds to bases 110-130 of the pareful of the primers primer and corresponds to bases 110-130 of the pareful of the primers primers and corresponds to bases 10-130 of the pareful of the primers and primers and corresponds to bases 10-130 of the pareful of the primers primers and primers and corresponds to bases 10-130 of the pareful of the primers primers and primers and corresponds to bases 10-130 of the pareful of the primers primers and primers and corresponds to bases 10-130 of the pareful of the primers primers and primers and corresponds to bases 10-130 of the pareful of the primers primers and primers and corresponds to bases 10-130 of the pareful of the primers primers and primers and corresponds to bases 10-130 of the pareful of the primers primers and primers and corresponds to bases 10-130 of the pareful of the primers primers and primers and corresponds to bases 10-130 of the pareful of the primers primers and primers and corresponds to bases 10-130 of the pareful of the primers primers and primers primers primers primers primers primers proving the primers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Promoting or suppressing corneal cell proliferation - usi hepatocyte growth factor or calcium ions resp., e.g. for corneal injury or for preserving corneal tissue prior to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US5589451-A.
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Column 11-12; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transplantation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-076878/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wilson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TEXA ) UNIV
   21
BP; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92US-0947683.
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A;
3 C; 6 G; 3 T;
   0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          growth factor
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Ś Matches Query Match **Local** 1322 TATCCTTTCACTCTGCATGGT 1342 l Similarity 21; Conserv Conservative 1.5%; Score 21; pred. No. 0; Mismatches DB 1; 0 Length 21; Indels 0 Gaps 0

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밁
AAV05498 standard; DNA;
                                       21
21
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01-MAY-1998 AAV05498;

(first entry)

RESULT 7
AAVO5498/c
ID AAVO54
XX
AAC AAVO54
XX
DT 01-MAY
DT 01-MAY
XX
DE Downst
XX
Inhibi
KW Inhibi
KW Lepato
KW dry ey
KW dry ey
KW fibrob
XX
XX
XX
Synthe
OS Synthe Inhibition; corneal epithelial cell; differentiation; treatment; hepatocyte growth factor; HGF; keratinocyte growth factor; KGF; dry eye; keratoconjunctivitis sicca; PCR primer; receptor; fibroblast growth factor; FGF; ss. Synthetic Downstream primer for FGF receptor DNA.

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RAESULLT

AAX3489

ID AAA

AC AA

AC AA

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CO AA

CO AA

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AC 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JLT 8
34892/c
Immortalized human hair papila cells used for evaluation of hair growth agent - are prepared by transformation of human hair papil cells with gene with deleted replication initiation point
                                                                                                                                                                                                                                                                                                                                                                                                                                              SV40 viral Large T-antigen gene; deleted repli
hair growth stimulating agent; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polymer microcapsules. The HGI topically. The method is used keratoconjunctivitis sicca.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KGF are both used, the cells can be contacted with them sequentially or simultaneously. The HGF and/or KGF is in a timed release delivery system, especially comprising biodegradable polymer microcapsules. The HGF and/or KGF are administered
                                                                                                                                                                                                                                                    19-SEP-1997;
                                                                                                                                                                                                                                                                                                     06-APR-1999.
                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immortalized human hair papila cell; HPC; screening; hair growth; SV40 viral Large T-antigen gene; deleted replication initiation point;
                                                                                                                                                                                                 19-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                   JP11089565-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wilson
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21-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-MAR-1995;
                                                                                                                                                     (SHIS ) SHISEIDO CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Inhibition of corneal cell differentiation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-076459/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (TEXA ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-DEC-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US5703047-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present sequence was used in the development of a novel method the inhibition of corneal epithelial cell differentiation. The hod comprises contacting the cells with a hepatocyte growth tor (HGF) and/or keratinocyte growth factor (KGF). When HGF and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer used to amplify FGFR2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     factor and/or keratinocyte growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TATCCTTTCACTCTGCATGGT 1342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Columns 17-18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TEXAS SYSTEM
                                                                                                                                                                                                   97JP-0271927
                                                                                                                                                                                                                                                    97JP-0271927.
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92US-0947683.
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                                                                                                                                                       LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 C; 6 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       36pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for treating dry eye,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 γģ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    especially
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                       papila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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                                                                                                        Matches
                                                    Query Match
Best Local
                                                                                                                                    The specification describes the preparation of immortalized human hair papila cells (HPC). The method comprises transformation of HPC with an SV40 viral Large T-antigen gene with deleted replication initiation point. The immortalized HPC can be used in a screening method for a hair growth agent, by culture of immortalized HPC in the presence of a substance to be tested and observation of the growth of the immortalized HPC. HPC is also used in development of hair growth stimulating agents. The present sequence represents a PCR primer, which is used in the course of the invention.
                                                                                                     Sequence 20 BP; 1 A; 4 C; 4 G; 11 T; 0
                                                                                                                                                                                                                                                                                                           Example
     1470
                                    20;
                                                       Similarity
                                                                                                                                                                                                                                                                                                           2; Page 7; 23pp; Japanese
   AATGAAGAACACGACCAAGA 1489
                                      Conservative
                                                    100.0%;
                                                                     1.4%;
                                    0;
                                                                       Score 20;
                                                       Pred. No.
                                      Mismatches
                                                                                                        other;
                                                      DB 1;
22;
                                      0
                                                                   Length
                                      Indels
                                    0,
                                      Gaps
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20

AATGAAGAACACGACCAAGA 1

0

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RESULT 9
AAXO0032/c
ID AAXO0032;
XX
AC AAXO0032;
XX
AC AAXO0032;
XX
DT 16-MAR-1999 (first entry)
XX
DE FGFR-2 (bek) PCR antisense
XX
Neuroepithelial stem cell;
XX
Neuroepithelial
XX
Neuroepithelial
XX
Neuroepithelial
XX
Neuroepithelial
AX
Neu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Neuroepithelial stem cell; lineage restricted intermediate procursor; oligodendrocyte; astrocyte; self-renewal; neuron; differentiation; neural creat cell; fibroblast growth factor; FGF, FGFF; receptor; CNS; central nervous system; glial cell; PCR primer; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98US-0073881.
97US-0852744.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98WO-US09630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antisense
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FOUND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rao MS
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The present invention describes an isolated, pure population of mammalian neuroepithelial stem cells, which are capable of self-renewal in adherent feeder-cell-independent (ARCI) culture medium and differentiation to central nervous system (CNS) neuronal or glial cells and to neuronal crest stem cells. Also described is an isolated population of mammalian CNS glial-restricted precursor (GRP) cells which can self-renew in the APCI culture medium and can differentiate to CNS glial cells but not to CNS neuronal cells. The stem cells can be used to generate a population of mammalian motor neurons by incubating the stem cells in a medium promoting cell proliferation and neuronal

Mammalian neuroepithelial stem cells and glial restricted precursor - can self renew and differentiate into central nervous system cells, used for generating various types of cells

78pp; English

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RESULT 10
AAT84432
ID AAT84432
AC AAT84
XX AAT84
AC AAT84
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Best Local :
                               claimed method for identifying the presence or absence of a deletion of the KIT gene sequence in pig genomic DNA. Other claimed primers (see AAT84420-27) are used to detect a duplication of the KIT gene. The 3 alleles for coat colour (I, inhibition of coat colour; I(p), patch; and i, development of colour) can be differentiated on the basis of duplication/deletion in the KIT gene. This allows breeding of pigs with the desired, usually white, coat colour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      medium lacking chick embryo extract. The stem cells can also produce neural crest stem cells by inducing the cells to differentiate in vitro. The inducing step comprises replating the cells on a laminin-coated substrate and preferably withdrawing a mitogen (preferably fibroblast growth factor; FGF) and chick embryo extract. Inducing can also comprise adding a dorsalizing agent to the cells, preferably a bone morphogenetic protein (BMP) such as BMP-2, -4 or -7. The stem cells can be used to produce cells of the peripheral nervous system, by inducing the stem cells to differentiate in vitro to neural crest stem cells, and inducing these cells to differentiate. AAX00029 to AAX00054 represent PCR primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Determn. of coat colour genotype in pigs by analysis of the KIT gene - for duplication or deletions, or analysis of KIT protein, used to establish breeding programmes for pigs of selected colour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-DEC-1995;
27-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KIT gene; pig; coat colour; pigmentation; primer; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                     Claim 39; Page 43; 49pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1997-145712/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Andersson L,
Wales R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-FEB-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9705278-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-NOV-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (DALG-) DALGETY PLC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KIT gene primer KITDEL1-FOR
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                                                                                                                                                                                                                                                                                                                                      pairs KITDEL1-FOR (AAT84432) and KITDEL1-REV (AAT84433)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20;
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95GB-0015385.
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                                                                                                                                                                                                                                                                                                                                                                                                     English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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22;
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                                                                            A method has been developed for: (a) differentiating animals and animal CC products according to breed origin; (b) determining or testing the breed CC origin of a product; or (c) validating an animal product. The method CC comprises analysing a sample of the product for the allele(s) of at C cleast one breed-determinant (BD) gene. The present invention also CC describes: (1) methods for determining the coat colour genotype of a pig CC by determining; (i) the allele(s) of the alpha melanocyte-stimulating CC hormone receptor (alpha-MSHR) gene; (ii) the amino acid sequence of an CC alpha-MSHR protein at positions associated with coat colour, or the size CC of the protein; (iii) detecting which microsatellites (or other linked CC marker alleles), linked to the alpha-MSHR gene, or particular alleles of it, are present; and (iv) analysing nucleic acid to determine if the KIT CC gene carries a polymorphism associated with the Belt genotype. The CC main method of the invention is applied to samples from fish, birds and CC marmals, especially pigs. Particular applications are confirming stated CC origin of meats; in quality control; for maintaining stock purity, and CC in breeding programmes (to confirm particular crosses). The method CC invention of meats; The process can be made quantitative. The present sequence represents a KIT gene PCR primer from the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Differentiating products from different animal breeds - by the analysis of alleles of breed-determinant genes, at the nucleic or protein level
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Porcine; wild boar; meishan; pietrain; large white; hampshire; differentiation; breed origin; alpha-MSHR; coat colour; stock problem melanocyte-stimulating hormone receptor; KIT; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV80714;
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97GB-0011214.
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Pred. No. 4:
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                                                                                                                                                                                                                         The invention provides a composition comprising one or more antisense oligonucleotides directed against vascular endothelial growth factor (VEGF) where the antisense oligonucleotides inhibits proliferation of cells exhibiting autocrine VEGF activity at an IC 5 0 concentration of between 0.5-2.5 micro Ma. The antisense oligonucleotides may be directed against VEGF for inhibiting cancer cell proliferation and angiogenesis. Preferably the oligonucleotide AAH23032 (a modified version of AAH22984) is used and may be utilized to treat Kaposi's sarcoma, ovarian cancer, prostrate cancer, pancreatic cancer or melanoma. Sequences AAH23012-023 represent gene-specific primers used in RT-PCR amplification of VEGF
                                                                                                                                                                                                                                                                                                                                                                                                   Novel antisense oligonucleotides useful for inhibiting vascular endothelial growth factor expression, angiogenesis and for treating cancer, e.g., Kaposi's sarcoma, ovarian cancer and prostrate cancer
            ABX90509 standard; DNA; 23
                                                                                                                                                                                       Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense therapy; RT-PCR; primer; VEGFR-1; ss.
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                                                                                                                                                                                                                                                                                                                                                                            Example 12; Page 56; 105pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vascular endothelial growth factor; VEGF; antisense; angiogenesis;
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                                                                                                         2098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2351 TGTGGGAGATCTTCACTTTAGG 2372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  _
                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                       CAGCTGGCCAGAGGCATGGAGT 2119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TGTGGGAGCTCTTCTCTTTAGG 22
                                                                                 CAAGTGGCCAGAGGCATGGAGT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Masood R;
                                                                                                                                1.3%;
ilarity 90.9%;
Conservative
                                                                                                                                                                                       BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               specific forward primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                       ?
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kaposi's sarcoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.3%;
                                                                                                                                                                                       4
                                                                                                                                                                                       C; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23 BP
              ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>..</u>
                                                                                                                                                                                       G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 18:8;
Pred. No. 41;
                                                                                                                                             Score 18.8;
Pred. No. 43
                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer; melanoma; cytostatic;
                                                                                                                                                            DΒ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               В
                                                                                                                                                           1;
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                                                                                                                                   <u>ب</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 22;
                                                                                                                                                           Length 23;
                                                                                                                                  Indels
                                                                                                                                  <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                  Gaps
                                                                                                                                   0
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RESULT 14
AAA95390
ID AAA95
XX AAA95
AC AAA95
XX 12-FE
XX 12-FE
XX Rat;
XW Rat;
XW Parki
XX Parku
XX Rat;
XW Parku
XX Rat;
XW Parku
XX Rat;
XW Parku
XX Rat;
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                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-JAN-1997;
19-JAN-2001;
30-JAN-1998;
19-JAN-2000;
                                                                                                                                                                                                                                                                                                                                                                                     cellular proliferation and preventing angiogenesis. The present sequence is a reverse transcriptase (RT)-PCR primer for a VEGF or related gene, used to clone the coding region for expression in tumour cell lines. The cell lines were used to test prospective antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a composition comprising an antisense oligonucleotide directed against vascular endothelial growth factor (VEGF). The antisense oligonucleotide is useful for preparing a composition treating cancer, neoplastic proliferation, abnormal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense; ss; PCR; VEGF; vascular endothelial growth factor; human; cancer; angiogenesis; neoplastic proliferation; cellular proliferation; primer; RT-PCR; reverse transcriptase PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gill PS,
                        Rattus norvegicus
                                                 Rat; Nurr1; tyrosine hydroxylase; catecholamine-related Parkinson's disease; manic depression; schizophrenia; Po
                                                                                                                      12-FEB-2001
                                                                                                                                                                         AAA95390 standard;
                                                                                                                                                                                                                                                                                                                                                 Sequence 23 BP; 6 A; 4 C; 9 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 12; Page 19; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       composition for treating cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New composition comprising an antisense oligonucleotide directed against vascular endothelial growth factor, useful for preparing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-246674/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GILL/)
(MASO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-MAR-2001; 2001US-0805761.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-NOV-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US2002165174-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human VEGFR-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAY-2003
                                                                                                                                                                                                                                                                                                           Local
                                                                                          FGFR coding sequence PCR primer #1.
                                                                                                                                                                                                                                                                   2098 CAGCTGGCCAGAGGCATGGAGT 2119
                                                                                                                                                                                                                                          ш
                                                                                                                                                                                                                                                                                            . Similarity
20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GILL PS.
MASOOD R.
                                                                                                                                                                                                                                          CAAGTGGCCAGAGGCATGGAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Masood
                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001WO-US00019.
98US-0016541.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000US-0487023
                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RT-PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97US-037004P.
)01WO-US00019.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      æ
                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                         90.9%;
                                                                                                                                                                                                                                                                                                                      1.3%;
                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     #1
                                                                                                                                                                                                                                                                                                          Score 18.8;
Pred. No. 43;
                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                             2
                                                                                                                                                                                                                                                                                                                      Length 23;
                                                                                                                                                                                                                                                                                               Indels
                                                   PCR primer;
                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                             Gaps
                                                      88
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WO200058451-A1

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RESULT 15
AAV60732/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The Nurr1 gene and protein can be used in the treatment of catecholamine-related diseases such as Parkinson's disease, manic catecholamine-related diseases such as Parkinson's disease, manic catecholamine related they can additionable used to induce tyrosine depression and schizophrenia. They can always use related hydroxylase expression and identify tyrosine hydroxylase related deficiencies, which are linked to the same diseases. The present sequence is a PCR primer used in a method to differentiate adult neural progenitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes the rat Nurrl coding and protein sequences. The Nurrl protein is involved in the induction of tyrosine hydroxylase expression in adult rat-derived hippocampal progenitor cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cell comprising exogenous nucleic acid inducing tyrosine hydroxylase expression useful for treating catecholamine-related diseases such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                             PCR primer; amplification; yeast; UAS; upstream activating sequence; UAS; transcription terminator; cell cycle; Upstream Activation Sequence; UAS; promoter; phosphorylation; cyclin; cyclin-dependent kinase; CDK; vector; cyclin kinase inhibitor; CKI; growth; wound healing; cancer therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sakurada K,
WPI; 1998-251302/22
                                                                              27-NOV-1996;
16-OCT-1996;
                                                                                                                      16-OCT-1997;
                                                                                                                                                                            WO9816660-A1
                                                                                                                                                                                                         OMO
                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                   Primer #2 for human CDK2 codons 1-149.
                                                                                                                                                                                                                                                                                                                                             08-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                        AAV60732;
                                                                                                                                                                                                                                                                                                                                                                                                 AAV60732 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 20; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Parkinson's disease, manic depression and schizophrenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SALK ) SALK INST BIOLOGICAL STUDIES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-MAR-2000; 2000WO-US07544.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-OCT-2000
                                                    (BITT-) BITTECH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1870 TCAGAGATGAGATGAA 1889
                                                                                                                                                                                                      sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      _
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Palmer T,
                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0277078
                                                                             96US-0031968.
96US-0029127.
                                                                                                                        97WO-US18608.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; 1 C; 7
                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.3%;
90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                   20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gage FH;
                                                                                                                                                                                                                                                                                                                                                                                                   BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 4 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 18.6;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     멂
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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used to screen for compounds that affect mammalian cell cycle regulatory or proteins. The method comprises administering a compound to a cell line, which contains a reporter gene linked to an Upstream Activation Sequence (CAS) and a promoter, where the UAS binds a transcription control factor (TCF) which is regulated through cyclin/cyclin-dependent kinase (CDK) phosphorylation. Also included in the construct is an effector gene providing a gene product to permit normal cyclin/CDK regulation of the reporter gene is then analysed in the cell line, TCF. Expression of the reporter gene is then analysed in the cell line, thereby determining whether the compound affects the normal regulation. The method can be used to identify inhibitors and activators of cyclins, CDKs, cyclin/CDK complexes, cyclin kinase inhibitors (CRIs), and cyclin/CDK/CKI complexes. The identified agents can be used for stimulating growth of cells (as in wound healing), or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Screening for agents that effect cell cycle regulatory proteins using a cell line that expresses a reporter gene in response to regulation through phosphorylation by a cyclin/CDK system
Sequence 20 BP; 4 A; 5 C; 5 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Primers AAV60731-V60732 were used to PCR amplify codons 1-149 of the human cyclin-dependent kinase 2 (hCDK2) gene. The amplified product were to generate a fusion protein comprising part of the hCDK2 sequentianked to codons 154-302 of the yeast PHO85 gene. The fusion protein
                                                                                                                regulating excessive cell growth and division (as in cancer therapy).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The fusion protein is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the hCDK2 sequence
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Matches
                                          Query Match
Best Local
                2201
20 CAGACTTTGGACTAGCCAGA 1
                                  19;
                                           Similarity
         CAGACTTTGGACTCGCCAGA 2220
                                  Conservative
                                          1.3%;
                                  0
                                           Score 18.4;
Pred. No. 4
                                   Mismatches
                                                   DB 1;
                                  1;
                                                 Length
                                   Indels
                                  0
                                   Gaps
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20;

0

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DRX PPI
                                                                                                                                                                                                            AAQ33179 standard; DNA; 24
                                                                                                                                           neoplasm; cancer; oncogene; tumour; growth; detection; diagnosis;
prognosis; treatment; sporadic colorectal carcinomas; ss.
                                                                                                                                                                                               AAQ33179;
                                                                                                      06-AUG-1992.
                                                                                                                   WO9213103-A1.
                                                                                                                                                               PCR primer #3
                                                                                                                                                                            28-JAN-1993
                                                                                                                                                                                  25-MAR-2003
                                                                                                                                                                            (updated)
(first entry)
                                                                                                                                                               to identify subtle DP2.5 mutations.
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(CANC-) (ICIL) (UYJO) (UTAH)

IMPERIAL CHEM IND PLC UNIV JOHNS HOPKINS.

CANCER INST

16-JAN-1991; 08-AUG-1991;

91GB-0000963. 91US-0741940. 92WO-US00376

16-JAN-1992;

WPI; 1992-284685/34 White RL, Nakamura Y;

Albertsen

Ή

Vogelstein B, Carlson ML,

Anand Groden

ĮĮ,

Hedge PJ, Joslyn G,

Markham AF; Thliveris

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UNIV UTAH.

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RESULT 17
AAT95598/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 20; Conserv
         Albertsen H,
Joslyn G, Kir
Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene. It was used with AAQ33180. To obtain DNA sequence adjacent to the exons of the gene, sequencing substrate was obtained by inverse PCR amplification of DNAs from two YACS 310D8 and 183H12 that span the deletions. Ligation at low concentration cyclized the restriction enzyme digested YAC DNAs. Oligonucleotides with sequencing tails designed in inverse orientation at intervals along the cDNAs primed PCR amplification from the cyclised templates. Comparison of these DNA sequences with the cDNA sequences placed exon boundaries at the divergence points. NOTE: The sequence as given in the specification is barely legible. See also AAQ33158-253.

(Updated on 25-MAR-2003 to correct PN field.)
                                                              (NICA-)
(UYJO)
(UTAH)
(ZENE)
                                                                                                                                                                                                                                                                                                                         Human; adenomatous Polyposis coli; APC; diagnosis; prognosis; neoplastic tissue; tumour tissue; tumour repressor; mutation; sporadic colorectal cancer; detection; PCR primer; DP2.5; SSCP; single stranded conformation polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection of somatic and germ-line alterations of human APC - used to diagnose, treat and study familial adenomatosus polyposis and sporadic colorectal cancer
                                                                                                                                                     16-JAN-1991;
16-JAN-1991;
                                                                                                                                                                                16-JAN-1991;
                                                                                                                                                                                                          12-AUG-1994;
                                                                                                                                                                                                                                    15-JUL-1997.
                                                                                                                                                                                                                                                            US5648212-A.
                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                            Primer for SSCP analysis of DP2.5 (APC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT95598;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT95598 standard; DNA; 24 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 24 BP; 10 A; 3 C; 4 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 8; Table 3; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2645 CTTCAGGAGATGATTCTGTTTTT
                                                                                                                                                                                                                                                                                      sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer was used to detect subtle mutations in the DP2.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23
                                                                          JAPANESE FOUND CANCER UNIV JOHNS HOPKINS. UNIV UTAH.
                                                              ZENECA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTTCAAGACATGATTCTGTATTT
          Kinzler K, M
B, White RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
            Anand R, Carry K, Markham A,
                                                                                                                                                                                                                                                                                                                                                                                                                    (updated)
(first en
                                                                                                                           91GB-0000962.
91GB-0000963.
91GB-0000974.
91GB-0000975.
91US-0741940.
                                                             LTD.
                                                                                                                                                                                                          94US-0289548
                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.3%;
                                    Carlson M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 18.2;
Pred. No. 58
                                                                                                    RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2667
                       Groden J,
Nakamura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1;
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•
                     Hedge PJ;
Thliveris A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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RESULT 18
AAV56500/c
ID SAV56
XX AAV56
XX AAV56
XX AAV56
XX AAV56
XX Pamil
XW Famil
XW Chrom
XX Famil
XW FaD;
XX Chrom
XX FAD;
XX FAD;
XX Chrom
XX FAD;
XX FAD;
XX Chrom
XX Chrom
XX FAD;
XX Chrom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                문
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                    16-JAN-1991;
16-JAN-1991;
08-AUG-1991;
12-AUG-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is a primer for the SSCP analysis of DP2.5 (APC), which was used in the development of a novel method of diagnosing or prognosing a human adenomatous Polyposis coli (APC) gene associated neoplastic tissue. The method comprises comparing APC gene coding sequences or mRNA in a non-neoplastic tissue, where a difference indicates an APC gene associated meoplasia of the tumour tissue. APC is a tumour repressor expressed in most normal tissues. APC mutations are found in familial adenomatous polyposis and sporadic colorectal cancer patients. The method enables mutations to be detected to provide an indication of predisposition to
                                         Albertsen H, Anand R, Carlson
Joslyn G, Kinzler K, Markham
Vogelstein B, White RL;
                                                                                                                                                (CANC-)
(UYJO )
(UTAH )
(ZENE )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Familial adenomatous polyposis coli; APC; tumour suppressor; therap
chromosome 5q21; tumourigenesis; retinoblastoma; colorectal tumour;
FAP; Gardner's Syndrome; GS; predisposition; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human DP2.5
WPI; 1998-427100/36
                                                                                                                                                                                                                                                                                                                                                                     16-JAN-1991;
16-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US5783666-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
23-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV56500;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV56500 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 BP; 10 A; 3 C; 4 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 8; Columns 31-32; 140pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cancer diagnosis - by detecting mutation(s) in adenomatous polyposis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-372053/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (Updated on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2645 CTTCAGGAGATGATTCTGTTTTT 2667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23
                                                                                                                                             UNIV JOHNS HOPKINS
UNIV UTAH.
ZENECA PHARM.
                                                                                                                                                                                                                          CANCER INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APC primer #23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (updated)
(first entry)
                                                                                                                                                                                                                                                                    91GB-0000974.
91GB-0000975.
91US-0741940.
94US-0289548.
                                                                                                                                                                                                                                                                                                                                                                     91GB-0000962
                                                                                                                                                                                                                                                                                                                                                                                                                                                   95US-0452655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.3%;
                                                                        Carlson M,
Markham AF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 18.2;
Pred. No. 58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                        Groden J, l
Nakamura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ω
                                                                          Hedge PJ;
, Thliveris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24;
                                                                          P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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RESULT 19
ANT962313/c
ID ANT962
XX ANT962
XX ANT962
XX 25-MAR
DT 08-APR
DT 08-APR
XX Human;
KW Human;
KW SSCP;
XX Synthe
OS Homo s
XX US5691
XX US5691
XX US5691
XX US5691
XX I6-JAN
PR 16-JAN
PR 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                (CANC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; adenomatous Polyposis coli; APC; diagnosis; prognosis; neoplastic tissue; tumour tissue; tumour repressor; mutation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      adenomatous polyposis (FAP) and Gardner's Syndrome (GS). The protes be used in therapy to replace lack of native functional protein and nucleic acids can be used for gene therapy. The nucleic acids that encode them can also be used as probes and primers in detection of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for the protein is present on human chromosome 5q21 and is also referred to as adenomatous polyposis coli gene. It is a tumour suppressor gene, and mutations in this gene have been associated with tumourigenesis in retinoblastoma and colorectal tumours, and especially familial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    neoplastic tissue; t
sporadic colorectal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Adenomatous polyposis coli protein cancers associated with mutation(s)
                            WPI; 1998-017712/02
                                                                        Albertsen H, Anand R, Carlson M,
Joslyn G, Kinzler K, Markham AF,
Vogelstein B, White RL;
                                                                                                                                                                                                                                                                                                                                16-JAN-1991;
08-AUG-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                16-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US5691454-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer for SSCP analysis of DP2.5 (APC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT96213 standard; DNA; 24 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24 BP; 10 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancers and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          adenomatous polyposis coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV56477-V56581 are primers used in the isolation of a human familial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example
                                                                                                                                                                                                                                                                                                                                                                               L6-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                         .6-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2645 CTTCAGGAGATGATTCTGTTTTT 2667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
                                                                                                                                                                                                   IMPERIAL CHEM IND PLC UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                       CANCER INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8; Column 31-32; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTTCAAGACATGATTCTGTATTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          predisposition to it. 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                 UTAH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       stranded conformation polymorphism; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (updated)
(first entry)
                                                                                                                                                                                                                                                                                                      91GB-0000975.
91US-0741940.
94US-0289548.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95US-0452654.
                                                                                                                                                                                                                                                                                                                                                                               91GB-0000974.
                                                                                                                                                                                                                                                                                                                                                                                                                                91GB-0000962.
                                                                                                                                                                                                                                                                                                                                                                                                         91GB-0000963.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 C; 4 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (APC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    detection; PCR primer; DP2.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 18.2;
Pred. No. 58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein from clone DP2.5. The gene

    useful in the treatment
    on human chromosome 5q21

                                                                                                  Groden J,
Nakamura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>س</u>
                                                                                                                             Hedge PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 24;
                                                                                                Thliveris A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The protein can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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RESULT 20
AAA93502/c
ID AAA935
XX AAA935
XX AAA935
XX Human
XX APC ge
KW Famili
KW Sporad
KW Sporad
KW Geneti
XX Homo s
XX US6114
XX I6-JAA
PR 16-JAA
PR 16-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                          Carlson M, G
Albertsen H,
Hedge PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diagnosing or prognosing a human adenomatous Polyposis coli (APC) gene associated neoplastic tissue. The method comprises comparing APC gene coding sequences or mRNA in a tumour tissue, to APC gene coding sequences or mRNA in a non-neoplastic tissue, where a difference indicates an APC gene associated neoplasia of the tumour tissue. APC is a tumour repressor expressed in most normal tissues. APC mutations are found in familial adenomatous polyposis and sporadic colorectal carger patients. The method enables mutations
                                                                                                                                                                                                                                                                                                                                  16-JAN-1991;
16-JAN-1991;
16-JAN-1991;
08-AUG-1991;
                               WPI; 2000-565003/52
                                                                                                                                                                                                         (ICIL )
(UYJO )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APC gene; Adenomatous Polyposis Coli gene; human; chromosome 5q21; familial adenomatous polyposis; FAP locus; Gardner's syndrome; GS; sporadic tumour; adenoma; carcinoma; cancer; lung; breast; colon; lbladder; liver; sarcoma; stomach; prostate; leukaemia; lymphoma; tumour suppressor; anti-APC antibody; detection; diagnosis; progno; genetic predisposition; drug screening; DP2.5; exon; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA93502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 BP; 10 A; 3 C; 4 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human APC (DP2.5) gene exon 2 PCR primer 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to be detected to provide an indication of predisposition to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 8; Columns 25-26; 107pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antibodies to normal and mutant adenomatous polyposis coli proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                          16-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US6114124-A.
                                                                                                                                                                                                                                                                                                                12-AUG-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (Updated on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2645 CTTCAGGAGATGATTCTGTTTT 2667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      which was used
                                                                                                                                                                                                      IMPERIAL CHEM IND PLC
UNIV JOHNS HOPKINS.
UNIV UTAH.
                                                                                                                                                                                   CANCER INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for detecting genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTTCAAGACATGATTCTGTATTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003 to correct PR field.)
                                                                                                       Groden J,
White RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                94US-0289548
                                                                                                                                                                                                                                                                                                                                     91GB-0000975
91US-0741940
                                                                                                                                                                                                                                                                                                                                                                                          91GB-0000974
                                                                                                                                                                                                                                                                                                                                                                                                                  91GB-0000963
                                                                                                                                                                                                                                                                                                                                                                                                                                          91GB-0000962.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95US-0450582.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              is a
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                                                                                                    Joslyn G, Kin:
,, Thliveris A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24
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Pred. No. 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               predisposition
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                                                                                                                               Kinzler K, Markham AF,
                                                                                                       Nakamura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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                                                                                                       Vogelstein
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                                                                                                       Anand R;
tein B;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        prognosis;
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RESULT 21

ABS67171/c

ID ABS67171 standard; DNA; 24 BE

XX

AC ABS67171;

XX

DT 29-NOV-2002 (first entry)

XX

DT, SRP19, DP25 gene SSCP pr

XX

Adenomatous polyposis coli; i

KW Adenomatous polyposis coli; i

KW mutation detection; tumour; c

KW single stranded conformations

XX

BY

CONTROL OF CONTROL

XX

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Best Local S
Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Adenomatous polyposis coli; APC; human; neoplastic tissue; mutation detection; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 24
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          91GB-000962.
91GB-0000973.
91GB-0000974.
91GB-0000975.
91US-0741940.
94US-0289548.
                                                                                                                                                                                                                                            95US-0449731.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  conformational polymorphism; primer; ss
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Pred. No. 58
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RESULT 22
AAV60725/c
ID AAV607
XX
AC AAV607
XX
Primer
XX
PCR pr
KW PCR pr
KW PCR pr
KW PCR pr
KW Cyclin
XX
Synthe
OS Synthe
OS Homo B
XX
PD WO9816
XX
PD 23-APF
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PR 16-OCT
XX
PR 16-OCT
XX
PR 16-OCT
XX
PR 16-T-
XX
PR (BITT-
XX
XX
WPI; 1
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to a novel method to aid in the diagnosis or prognosis of a neoplastic tissue of a human. The method involves detecting somatic alteration of wild-type adenomateus polyposis coli) protein in a tumour tissue isolated from a human (the alteration indicating neoplasia of the tissue). The method of the invention is useful in diagnosis or prognosis of a neoplastic tissue of a human the method is useful in detection of genetic predisposition to cancer. The present sequence represents a DNA sequence used in the method of the invention of the inv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Albertsen H
Kinzler K,
                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer; amplification; yeast; UAS; upstream activating sequence; transcription terminator; cell cycle; Upstream Activation Sequence; promoter; phosphorylation; cyclin; cyclin-dependent kinase; CDK; veccyclin kinase inhibitor; CKI; growth; wound healing; cancer therapy;
                                                                                                                                                            27-NOV-1996;
16-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-DEC-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Method to aid in the diagnosis/prognosis of neoplastic tissues in humans, by detecting somatic alteration of wild-type APC protein in tumor tissue isolated from human, the alteration indicating neoplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYJO)
(UTAH)
(NICA-)
(ZENE)
                                                                                                                                                                                                                                16-OCT-1997;
                                                                                                                                                                                                                                                                                                                        WO9816660-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAV60725 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24 BP; 10
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                                                                                                               (BITT-) BITTECH
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ZENECA LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anand
                                                                                                                                                            96US-0031968
96US-0029127
                                                                                                                                                                                                                                97WO-US18608
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CDK2 codons 1-151.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 18.2; u
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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, Thliveris A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
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lstein B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            CDK; vector;
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WPI; 1998-251302/22

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RESULT 23
AAZ18094/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The method can be used to identify inhibitors and activators of mammalian cell cycle regulatory proteins, especially inhibitors and activators of cyclins, CDKs, cyclin/CDK complexes, cyclin kinase inhibitors (CKIs), and cyclin/CDK/CKI complexes. The identified agents can be used for stimulating growth of cells (as in wound healing), or regulating excessive cell growth and division (as in cancer therapy).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primers AAV60724-V60725 were used to PCR amplify codons 1-151 of the human cyclin-dependent kinase 2 (hCDX2). The amplified product was used to generate a fusion protein comprising part of the hCDX2 sequence linked to codons 155-302 of the yeast PHO85 gene. The fusion protein is used to screen for compounds that affect mammalian cell cycle regulatory protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              proteins. The method comprises administering a compound to a cell line, which contains a reporter gene linked to an Upstream Activation Sequence (UAS) and a promoter, where the UAS binds a transcription control factor (TCF) which is regulated through cyclin/cyclin-dependent kinase (CDK) phosphorylation. Also included in the construct is an effector gene providing a gene product to permit normal cyclin/CDK regulation of the reporter gene is then analysed in the cell line, thereby determining whether the compound affects the normal regulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Screening for agents that effect cell cycle regulatory proteins using a cell line that expresses a reporter gene in response to regulation through phosphorylation by a cyclin/CDK system
Identifying and characterizing cells by comparing the pattern gene expression in a selected gene family
                                                                                                                                                                                                                                                                                                                                                                                                        Genetic proximity; gene expression; cell characterisation; homeobox gene, genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR;
                                                  P-PSDB;
                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PTK 2 gene specific primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21
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                                                    WPI; 1999-419113/35.
P-PSDB; AAY14629.
                                                                                                                                                                     16-OCT-1998;
29-DEC-1997;
                                                                                                                                                                                                                      28-DEC-1998;
                                                                                                                                                                                                                                                       08-JUL-1999.
                                                                                                                                                                                                                                                                                         WO9934016-A2
                                                                                                                                                                                                                                                                                                                                                                         primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                       kinase gene; protein phosphatase; P450; steroid receptor; cadherin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ18094 standard; DNA; 21 BP
                                                                                                                                      (GENE-) GENENA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2203 GACTITGGACTCGCCAGAGAT 2223
                                                                                                                                                                                                                                                                                                                        sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                      LTD.
                                                                                                                                                                     98IL-0126627
97IL-0122793
                                                                                                                                                                                                                      98WO-IL00625.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; 6 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 17.8; D
Pred. No. 55;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 21;
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                   of
f
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                                                                                                                                                                                                                                              AAZ18102/c
                                                                                                                                                                                                                                                    RESULT 24
                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                  Identifying and gene expression
                                                                                                                                                                    Genetic proximity; gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinse gene; protein phosphatase; P450; steroid receptor; cadherin;
                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 7 A;
    Claim 4; Page 42; 102pp; English
                                     WPI; 1999-419113/35.
P-PSDB; AAY14637.
                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                          AAZ18102
                                                                                                                                                                                                                                       AAZ18102 standard; DNA;
                                                         Vider B;
                                                                                   16-OCT-1998;
29-DEC-1997;
                                                                                                                                                                primer; ss.
                                                                                                                                                                                                             11-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 42; 102pp; English
                                                                                                       28-DEC-1998;
                                                                                                                                 WO9934016-A2
                                                                                                                                             Homo sapiens
                                                                                                                                                                                                PTK 6 gene specific primer.
                                                                      (GENE-) GENENA
                                                                                                                                                                                                                                                                                                  19;
                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                               AGTGATGTCTGGTCCTTCGGG 2343
                                                                                                                                                                                                                                                                         AGTGATGTCTGGTCCTATGGG
                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                             (first entry)
                                                                      Q.I.7
                                                                                   98IL-0126627
97IL-0122793
                                                                                                       98WO-IL00625
                  characterizing cells by comparing the pattern in a selected gene family
                                                                                                                                                                                                                                                                                                       1.3%;
                                                                                                                                                                                                                                                                                                                           8 C;
                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                                                                                                                           3 G; 3 T; 0
                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                        Score 17.8;
Pred. No. 5
                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                            other;
                                                                                                                                                                                                                                                                                                          55
                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                  2
                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                               21;
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RESULT 25
AAZ18110/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transformed. They can be used for detecting a selected genetic defect in an individual, e.g. a fetus. They can also be used for determining the effect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired property. The method uses reverse transcriptase polymerase chain reaction (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences ARZ17803-218342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, protein phosphatase genes, P450 enzyme genes, steroid receptor superfamily genes or cadherin superfamily genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genetic proximity; gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin;
                                                                                                                                           WPI; 1999-419113/35.
P-PSDB; AAY14645.
                                                                                                                                                                                                                                                                                                                                        16-OCT-1998;
29-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is
                                                  Identifying and characterizing cells by comparing the pattern gene expression in a selected gene family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-JUL-1999
                                                                                                                                                                                                                                                                                      (GENE-) GENENA
                                                                                                                                                                                                                                                                                                                                                                                                                              28-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9934016-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PTK 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-OCT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ18110 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 7 A; 8 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention provides a new method for identifying and characterising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sapiens
  4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGTGATGTCTGGTCCTTCGGG 2343
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     specific
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
42;
                                                                                                                                                                                                                                                                                                                                        98IL-0126627.
97IL-0122793.
                                                                                                                                                                                                                                                                                                                                                                                                                              98WO-IL00625
102pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer.
English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 17.8;
Pred. No. 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21,
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The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the secon

cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and the second a second cell comprises: (a) obtaining the first cell and the second

Claim 4; Page 43; 102pp; English.

proximity index using a specified

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RESULT 26
AAZI8118/c
ID AAZI81
XX AAZI81
XX AAZI81
XX Geneti
XX Primer
XX W09934
XX W09934
XX O8-JUL
XX 28-DEC
XX O8-JUL
XX 16-OCT
PR 29-DEC
XX (GENE-
XX Vider
XX Vider
XX Geneti
PT Gene e
XX The in
CC cells.
CC of exp
CC of exp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genetic proximity; gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin;
                                                                                                                                                                                                              P-PSDB; AAY14653.
                                                                                                                                                                                                                                                                                                                                     16-OCT-1998;
29-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression. The gene family can be selected from a set of homeobox kinase genes, protein phosphatase genes, P450 enzyme genes, steroid receptor superfamily genes or cadherin superfamily genes.
                                                                                                                                                                                                                                WPI; 1999-419113/35.
                                                                                                                                                                                                                                                                 Vider B;
                                                                                                                                                                                                                                                                                                                                                                                        28-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                             08-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9934016-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PTK 14 gene specific primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ18118 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 7 A; 8 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cell; (b) determining in the first cell and the second cell the pattern
                                                                                                                                                                             [dentifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2323 AGTGATGTCTGGTCCTTCGGG 2343
                                                                                                                                                         tifying and characterizing cells by comparing the expression in a selected gene family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cal Similarity
19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                  LTD
                                                                                                                                                                                                                                                                                                                                   98IL-0126627.
97IL-0122793.
                                                                                                                                                                                                                                                                                                                                                                                          98WO-IL00625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 17.8;
Pred. No. 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2;
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                                                                                                                                                                           pattern
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RESULT 27
AAZ18216/c
ID
AAZ18216;
XX
AAZ18216;
XX
AZ18216;
XX
DE
T1-OCT-1999 (first entry)
XX
DE
Tyrosine kinase gene speci
XX
Genetic proximity; gene ex
XX
Genetic defect; reverse tx
XX
W genetic defect; reverse tx
XX
W primer; ss.
XX
OS
Homo sapiens.
XX
OS
Homo sapiens.
XX
V
PD
O8-JUL-1999;
XX
PD
O8-JUL-1999;
XX
PD
XX
V
PD
O8-JUL-1998; 98NO-IL0062
XX
PP
C1-OCT-1998; 98IL-012662
PR
29-DEC-1997; 97IL-012275
XX
C1-OCT-1998; 98IL-012662
PR
29-DEC-1997; 97IL-012275
XX
PD
OR
P-PSDB; AAY14750.
XX
Vider B;
XX
Vider B;
XX
Vider B;
XX
Claim 4; Page 48; 102pp; 1
XX
CC
The invention provides a 1
CC calls. The method for detection of genes in 1
CC calls. The method for detection of genes in 1
CC characterising cells, eaglectic status, whether in 1
CC transformed. They can be 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
  The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is transformed. They can be used for detecting a selected genetic defect in
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                                                                                                                                                                                                                                                                                                                       Identifying and characterizing cells by gene expression in a selected gene fami
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tyrosine kinase gene specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 7 A; 8 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptor superfamily genes or cadherin superfamily genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2323 AGTGATGTCTGGTCCTTCGGG 2343
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     proximity; gene expression; cell characterisation; homeobox gene; defect; reverse transcriptase polymerase chain reaction; RT-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein phosphatase; P450; steroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98IL-0126627.
97IL-0122793.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98WO-IL00625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21
                                                                                                                                                                                                                                                                      English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 17.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer 409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                         family
                                                                                                                                                                                                                                                                                                                                               comparing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             receptor; cadherin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 21,
                                                                                                                                                                                                                                                                                                                                                  pattern
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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Matches Query Match Best Local

Similarity

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Score 17.4; Pred. No. 56

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RESULT 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     an individual, e.g. a fetus. They can also be used for determining the effect of a selected treatment on a test cell. They can also be used fo obtaining cells capable of expressing an homeobox related desired property. The method uses reverse transcriptase polymerase chain reaction (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences AAZ17803-Z18342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene
                                                                                                    The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent k other than cell-cycle dependent kell-cycle dependent kell-cycle dependent kell-cycle pendent kell-cycle pen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-DEC-2000
                                                The ribozyme is resistant to endon efficient in restenosis treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ribozyme; hairpin; hammerhead; gene therapy; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cdk2 ribozyme binding site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA82640;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 6 A; 9 C; 4 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          kinase genes, protein phosphatase genes, P450 enzyme genes, steroid receptor superfamily genes or cadherin superfamily genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression. The gene family can be selected from a set of homeobox genes,
                                                                                                                                                                                                                                                                                                        Disclosure;
                                                                                                                                                                                                                                                                                                                                                            New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-412314/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tritz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200032765-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA82640 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (IMMU-) IMMUSOL INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity es 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2323 AGTGATGTCTGGTCCTTCGGG 2343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGCGATGTGTGGTCCTTCGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Welch PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88.
                                                                                                                                                                                                                                                                                                     Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                is resistant to endonuclease activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98US-0110954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US28772
                                                                                                                                                                                                                                                                                                     49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90.5%;
                                                                                                                                                                                                                                                                                                  109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Barber JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 17.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Robbins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         55;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                      and hence is
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
recognition site; target; ribozyme binding site; eye disease; vulnera proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinase clbr. PCNA and Cyclin B1.

Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells.
                                                                                                 Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19
                                                                                                                                                              Cell-cycle dependent kinase cdk2 ribozyme binding site
                                                                                                                                                                                                                                                                                             AAH57802
                                                                                                                                                                                                                                                                                                                                                       AAH57802 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New hairpin and hammerhead ribozyme RNA encoding a cyclin or cell-cycle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tritz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200032765-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ribozyme; hairpin; hammerhead; gene therapy; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cdk2 ribozyme binding site #78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-DEC-2000
                                                                                                                                                                                                                               10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 49; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA encoding a cycl
PCNA and Cyclin B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-412314/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA82641;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA82641 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ribozyme is resistant to endonuclease activity and hence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ,_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGCAGACTTTGGACTCGCC 2217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGCAGACTTTGGACTAGCC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TAGCAGACTTTGGACTAGC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'n.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Welch PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          restenosis treatment.
                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0110954.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5 C; 5 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Barber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 17.4;
Pred. No. 56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Robbins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for inhibiting restenosis, of dependent kinase other than
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ۳.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                              SEQ ID NO:226.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cleaves
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CDK1
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RESULT 31 AAH57803

AAH57803

standard; DNA; 19

SX SX SX SX SX S

AAH57803; 10-SEP-2001

(first entry)

Cell-cycle dependent kinase cdk2 ribozyme binding site SEQ ID NO:227

Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme;

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                                                                                                                                                                                    The present invention describes a method for treating a proliferative CC skin or eye disease and scarring. The method involves administering a CT ribozyme (I) which cleaves RNA encoding a cytokine involved in CC inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle concleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, antisickling, CC dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, CC ophthalmological, vulnerary, keratolytic and virucide activities, and CC cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, CC squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing such as keloid, adhesion and hypertrophic or hypertrophic burn carring such as keloid, adhesion and hypertrophic or hypertrophic burn carring such as keloid, adhesion and hypertrophic burn the
                                                                    Matches
                                                                                      Query Match
Best Local
                                                                                                                                        Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating proliferative skin or eye diseases and scarring, ribozymes that cleave RNA encoding cytokines involved in matrix metalloproteinases, growth factors and cell-cycle of matrix metalloproteinases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               matrix metalloproteinase; growth factor; reductase; scarring; cytostat: antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma;
                                                                                                                                                                         exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 88; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       kinases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-300427/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Robbins JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200130362-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      basal cell carcinoma; seborrheic wart;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-OCT-2000; 2000WO-US29500.
                                                                                    Local
                                 2198
                                                                    18;
                                                                                    Similarity
                    TAGCAGACTTTGGACTCGC
                                                                                                                                        ВP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tritz R;
                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0161532.
                                                                                                                                    5 A;
                                                                                   1.2%;
                                                                                                                                        4 C;
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                                                                                                                                        G.
                                                                    0,
19
                                 2216
                                                                                      Pred.
                                                                                                   Score 17.4;
                                                                                                                                        5 T;
                                                                    Mismatches
                                                                                      No.
                                                                                                                                        0 other;
                                                                                      56,
                                                                                                     DB 1;
                                                                    1;
                                                                                                     Length 19;
                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammation,
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                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                     inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, antiseborrheic, antidiabetic, antisickling, cophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, cyuamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH57577 to AAH62099 represent sequences used in the
                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; NMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic antipsoriatic; dermatological; antiseborrheic; antidabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis, actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent
                  16-DEC-2002
                                                     ABS78766;
                                                                                    ABS78766 standard; DNA;
                                                                                                                                                                                                                                                                                                                   Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 88; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-OCT-2000; 2000WO-US29500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200130362-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sickle cell retinopathy; ss.
                                                                                                                                                                                                                2199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2001-300427/31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              eye
                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ent invention describes a method for treating a proliferative eye disease and scarring. The method involves administering a (I) which cleaves RNA encoding a cytokine involved in
                                                                                                                                                                                                                AGCAGACTTTGGACTCGCC 2217
                                                                                                                                                                             AGCAGACTTTGGACTAGCC 19
                                                                                                                                                                                                                                                                                                                     BP; 5
                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tritz R
                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0161532
                                                                                                                                                                                                                                                                                                                     A; 5
                                                                                                                                                                                                                                                              1.2%;
                                                                                                                                                                                                                                                                                                                     C; 5
                                                                                      22
                                                                                      ₽₽
                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                     G; 4 T; 0 other;
                                                                                                                                                                                                                                                                  Pred
                                                                                                                                                                                                                                                                  Score 17.4;
Pred. No. 56;
                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                  BB
                                                                                                                                                                                                                                                                                  ۲,
                                                                                                                                                                                                                                                                                  Length 19;
                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic;
                                                                                                                                                                                                                                                 Gaps
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23-MAR-2001; 26-MAR-2001; 27-MAR-2001;

22-MAR-2001

2001US-277833P. 2001US-278152P.

2001US-277791P.

28-MAR-2001; 30-MAR-2001; 02-APR-2001;

2001US-278894P. 2001US-279036P. 2001US-279344P. 2001US-280233P. 2001US-280802P.

02-MAY-2001;

2001US-288148P. 2001US-294821P. 2001US-335302P. 2001US-338375P.

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Hodgkin disease; Von Hippel-Lindau syndrome; Alzheimer's disease; stroke tuberous sclerosis; hypercalcaemia; Parkinson's disease; depression; Huntington's disease; cerebral palsy; epilepsy; Lesch-Nyhan syndrome; multiple sclerosis; attaia-telanglectasia; leukodystrophy; anxiety; pain; obesity; Crohn's disease; osteoporosis; inflammatory bowel disease; infertility; inflammatory bowel disease; attherosclerosis; hypertension; scleroderma; haemophilia; diabetes; pancreatitis; autoimmune disease; asthma, arthritis; immunodeficiency; HIV; viral infection; neurogenesis; bacterial infection; parasitic infection; graft-versus-host disease; cell differentiation; cell proliferation; haematopoiesis; wound healing;
20-MAR-2001;
21-MAR-2001;
                                                                                                                                                                                                                                                      08-MAR-2001;
09-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                           08-MAR-2002;
                                                                                                20-MAR-2001;
                                                                                                                                                            13-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200272770-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human NOVX forward primer Ag3477.
                                                                                                                                                                                              13-MAR-2001;
                                                                                                                                                                                                                            12-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NOVX; human disease; NOVX-associated disorder;
                                                                                                                                                                                                                                                                                                                                                           2002WO-US07283.
                                    2001US-277338P.
                                                               2001US-277239P.
2001US-277327P.
                                                                                                                                  2001US-276000P.
                                                                                                                                                                                                                            2001US-274849P.
2001US-275235P.
                                                                                                                                                                                                                                                                                           2001US-274281P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer; addiction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disease;
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CURAGEN CORP

Gangolli Spytek KA, Zerhusen ΕA, BD, Vernet CA, Patturajan M, Taupier RJ; Tchernev VT, Nn M, Gusev VY, Malyankar UM, Kekuda R, Pena CEA, Gerlach VL, Zhong Li L;

WPI; 2002-713508/77.

New NOVX polypeptides and polynucleotides, useful for preventing, diagnosing or treating NOVX-associated disorders, e.g. diabetes, multiple sclerosis, atherosclerosis, cancer, infections, osteopor Parkinson's disease osteoporosis

Example C; Page 200; 266pp; English.

The present invention relates to a new polypeptide (NOVX). The NOVX polypeptide, nucleic acid and antibody are useful in the manufacture of a medicament for treating a syndrome associated with a human disease, preferably a NOVX-associated disorder. The NOVX nucleic acids, polypeptides and antibodies are useful for treating, preventing or diagnosing diseases such as cancers, Hodgkin disease, preventing or diagnosing diseases such as cancers, Hodgkin disease, von Hippel-Lindau syndrome, Alzheimer's disease, stroke, tuberous sclerosis, hypercalcaemia, Parkinson's disease, Huntington's disease, cerebral palsy, epilepsy, Lesch-Nyhan syndrome, multiple sclerosis, ataxiatelangiectasia, leukodystrophies, addiction, anxiety, depression, pain, obesity, Crohn's disease, osteoporosis, inflammatory bowel disease, bowel disease, atherosclerosis, hypertension

999999999999998**%**

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Query Match
Best Local S
Matches 19
                   The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genetic
genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   infections, or graft-versus-host disease. The nucleic acids and polypeptides may also be used as targets for the identification of small molecules that modulate or inhibit e.g. neurogenesis, cell iderentiation, cell proliferation, hasmatopoiesis, wound healing and angiogenesis, in gene therapy, in generation of antibodies that bind immunospecifically to NOVX substances for use in therapeutic or diagnostic methods. The nucleic acids are used in the hybridisation probes, in chromosome mapping, tissue typing, preventive medicine, and pharmacogenomics. The present nucleic acid sequence represents a PCR primer that was used in the methods of the invention for amplification
                                                                                                                                                                                                                                          Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     scleroderma, haemophilia, diabetes, pancreatitis, autoimmune disease, asthma, arthritis, immunodeficiencies, HIV, viral, bacterial or parasinfections, or graft-versus-host disease. The nucleic acids and
                                                                                                                                                                                                                                                                                                                Identifying and characterizing cells by
                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-419113/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GENE-) GENENA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          kinase gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PTK 9 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-OCT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ18107 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human NOVX.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1886 TGAAGATGATTGGGAAACACAA 1907
                                                                                                                                                                                                                                                                                         expression in a selected
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        proximity; gene expression; cell characterisation; homeobox gene;
defect; reverse transcriptase polymerase chain reaction; RT-PCR;
gene; protein phosphatase; P450; steroid receptor; cadherin;
                                                                                                                                                                                                                                                                                                                                                               AAY14642
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                                                                                                                                                                                                                                             Page 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TGAACATGTTTGGAAAACACAA
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97IL-0122793.
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                                                                                                                                                                                                                                        102pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           primer
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                                                                                                                                                                                                                                          English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        묪
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Pred. No. 74;
0; Mismatches
                                                                                                                                                                                                                                                                                       gene
for detecting a selected genetic
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                                                                                                                                                                                                                                                                                            family
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                                                                                                                                                                                                                                                                                                             comparing
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                                                                                                                                                                                                                                                                                                                the
                                                                                                                                                                                                                                                                                                                pattern
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an individual, e.g. a fetus. They can also be used for determining the effect of a selected treatment on a test cell. They can also be used fo obtaining cells capable of expressing an homeobox related desired property. The method uses reverse transcriptase polymerase chain reaction (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences AAZ170803-Z18342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox gene kinase genes, protein phosphatase genes, P450 enzyme genes, steroid receptor superfamily genes or cadherin superfamily genes.
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AAZ18109;
                                                                     PTK 10 gene
                                                                         11-OCT-1999
                                                                                AAZ18109 standard; DNA;
                                                                     specific primer.
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Genetic proximity; gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin; primer; 88.

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08-JUL-1999
                       WO9934016-A2
                                                         Synthetic
                                            sapiens.
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16-OCT-1998; 29-DEC-1997; 28-DEC-1998; (GENE-) GENENA LID. 98IL-0126627. 97IL-0122793. 98WO-IL00625.

Vider B;

P-PSDB; AAY14644.

1999-419113/35.

Identifying and characterizing cells by comparing the gene expression in a selected gene family pattern o H

Claim 4; Page 42; 102pp; English.

The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is transformed. They can be used for detecting a selected genetic defect in an individual, e.g. a fetus. They can also be used for determining the effect of a selected treatment on a test cell. They can also be used for cells capable

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                 The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and the second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell the pattern comprises in a selected gene family; and (c) calculating a greatinity index using a specified formula. The methods can be used for comparative for the second cell; its genetic status, whether it carries a genetic defect, or whether it is comparative formed. They can be used for detecting a selected genetic defect in an individual, e.g. a fetus. They can also be used for determining the ceffect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired component. The method uses reverse transcriptase polymerase chain capable of family the pattern of gene expression in a selected formed for the method uses reverse transcriptase polymerase chain capable of family the pattern of gene expression in a selected formed family the pattern of gene expression in a selected formed family the pattern of gene expression in a selected formed family the pattern of gene expression in a selected formed family the pattern of gene expression in a selected formed family the pattern of gene expression in a selected formed family the pattern of gene expression in a selected family the pattern of gene expression in a selected family the pattern of gene expression in a selected family the pattern of gene expression in a selected family the pattern of gene expression in a selected family the pattern of gene expression in a selected family the pattern of gene expression in a selected family the pattern of gene expression in a selected family the pattern of gene expression in a selected family the pattern of gene expression in a selected family the pattern of gene expression in a selected family of general family of general fami
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P-PSDB; AAY14646.
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               The invention provides a new method for identifying and characterising CC cells. The method for determining the genetic proximity of a first cell cc and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a CC proximity index using a specified formula. The methods can be used for CC characterising cells, e.g. for determining the origin of a cell, its CC genetic status, whether it carries a genetic defect, or whether it is CC transformed. They can be used for detecting a selected genetic defect in CC an individual, e.g. a fetus. They can also be used for determining the ceffect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired concepts. The method uses reverse transcriptase polymerase chain CC property. The method uses reverse transcriptase polymerase chain CC selected gene family. Sequences AAZ17803-Z18342 represent primers that CC can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, transity and property can also be concept.
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97IL-0122793.
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RESULT 37
AAZ18189
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                                          The invention provides a new method for identifying and characterising CC cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its compared to the continuous cell in the continuous cell in the continuous cell in the continuous cell in the cell in the continuous cells. They can selected genetic defect in an individual, e.g. a fetus. They can also be used for determining the ceffect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired property. The method uses reverse transcriptase polymerase chain creation (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences AAZ17803-Z818342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, protein phosphatase genes, P450 enzyme genes, steroid
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                             receptor superfamily genes or cadherin superfamily genes
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29-DEC-1997;
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97IL-0122793.
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100.0%; Pred. No.
Live 0; Mismatches
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RESULT 38
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                           The invention provides a new method for identifying and characterising CC cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second CC and a second cell comprises: (a) obtaining the second cell the pattern of cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a CC proximity index using a specified formula. The methods can be used for CC characterising cells, e.g. for determining the origin of a cell, its CC genetic status, whether it carries a genetic defect, or whether it is CC transformed. They can be used for detecting a selected genetic defect in CC transformed. They can be used for determining the cell they can also be used for ceffect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired CC property. The method uses reverse transcriptase polymerase chain CC reaction (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences AAZ17803-Z18342 represent primers that CC can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, protein phosphatase genes, P450 enzyme genes, steroid CC receptor superfamily genes or cadherin superfamily genes.
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Best Local Similarity
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29-DEC-1997;
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97IL-0122793.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98IL-0126627.
97IL-0122793.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98WO-IL00625.
                                                                                                                             A; 3 C; 4 G; 6 T; 0 other;
                                                           1.2%; Score 17;
100.0%; Pred. No.
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                                         Mismatches
                                                             DB 1;
69;
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                                                                                 Length 20;
                                            Indels
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                                            Gaps
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RESULT 41
AAC62905/c
ID AAC629
XX
AC AAC629
XX
DT 06-FEB

standard;

DNA;

20

0

AAC62905 AAC62905

06-FEB-2001

(first entry)

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RESULT 40
RASZ93562/c
ID AAX293
XX AAX293
XX AAX293
XX JUST 10-JUN
XX JUST 10-JUN
XX JUST 10-JUN
XX Antise
KW JUST 10-JUN
XX Synthe
OS Homo s
XX WO9909
XX Synthe
OS Homo s
XX WO9909
XX 13-AUG
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XX 15-FEB
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XX WPI 13-AUG
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                                                                                                        Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                      The invention relates to antisense oligonucleotides that detect and modulate the expression of Jun N-terminal kinase (JNK) proteins. The oligonucleotides specifically hybridize to a nucleic acid encoding a NK1, JNK2 or JNK3 protein, and which modulate expression of these proteins. The oligonucleotides are useful for modulating JNK protein expression and cell cycle progression in cultured cells or animal cells. The oligonucleotides are also useful for modulating the phosphorylation of a protein that has been phosphorylated by a JNK protein, and the expression of a cellular protein that promotes one or more metastatic events. The oligonucleotides also form pharmaceutical compositions for treating animals with a hyperproliferative disease, and for inhibiting terms are animal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense oligonucleotide; Jun N-terminal kinase; JNK; hybridise; JNK1; JNK2; JNK3; cell cycle progression; phosphorylation; tumour; probe; hyperproliferative disease; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX29362 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides that detect and modulate the expression of Jun N-terminal kinase proteins - useful for treating hyperproliferative diseases and inhibiting tumor growth in animals, and for modulating protein phosphorylation by these proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-JUN-1999
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                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                      tumor growth in an animal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dean N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JNK3-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Page 102; 190pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-181060/15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                            2642
      17
                                         GTTCTTCAGGAGATGAT 2658
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GTTCTTCAGGAGATGAT 1
                                                                                                                                                                                                                                             BP; 6 A;
                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97US-0910629
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                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                             6 C; 3 G; 5 T;
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                                                                                                                                                    Score 17; DB 1;
Pred. No. 69;
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                                                                                                                           Mismatches
                                                                                                                                                                                                                                                0 other;
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                                                                                                                                                                                    Length 20
                                                                                                                           Indels
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                                                                                                                        Gaps
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RESULT 42
AAXOO049
ID AAXOO
XX Neuro
XX Neuro
XX Oligo
KW Oligo
KW Oligo
KW Oligo
XX AAXOO
XX Synth
OS Synth
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                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to antisense oligonucleotides (AAC62944-C63000, AAA96093-A96099 and AAA07993) that hybridise specifically to a nucleotide encoding a Jun N-terminal kinase (JNK2) protein, resulting in decrease of JNK2 expression and leading to induction of apoptosis. The present sequence is one such antisense oligonucleotide. The oligonucleotides of the present invention are useful for treating diseases or conditions with reduced apoptosis, e.g. cancer and cellular hyperproliferation. The oligonucleotides may also be used to increase the stimulation of apoptotic proteins, e.g. for treating Alzheimer's or Parkinson's disease, amylotrophic lateral sclerosis, retinitis, pigmentosa, epilepsy, myocardial infarction, stroke, obstructive jaundice, polycystic kidney and diabetes. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cellular hyperproliferation; Alzheimer's; Parkinson's disease; amylotrophic lateral sclerosis; retinitis; pigmentosa; epilepsy; myocardial infarction; stroke; obstructive jaundice; polycystic kidney; diabetes; Jun N-terminal kinase; ss.
                                           Neuroepithelial stem cell; lineage restricted intermediate precursor; oligodendrocyte; astrocyte; self-renewal; neuron; differentiation; neural crest cell; fibroblast growth factor; FGF; FGFR; receptor; CNS; central nervous system; glial cell; PCR primer; amplification; ss.
 Synthetic
Homo sapie
                                                                                                                      FGFR PCR sense primer.
                                                                                                                                                        16-MAR-1999
                                                                                                                                                                                                                 AAX00049 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 6 A; 6 C; 3 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel methods for reducing apoptosis comprising antisense oligonucleotides, useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-638427/61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mckay R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                             2642 GTTCTTCAGGAGATGAT 2658
                                                                                                                                                                                                                                                                                              17
                                                                                                                                                                                                                                                                                                                                                        l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5; Page 139; 160pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  may
                                                                                                                                                                                                                                                                                             GTTCTTCAGGAGATGAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dean NM,
                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Monia
                                                                                                                                                                                                                                                                                                                                                                     1.2%;
                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                                                                                                                                                      Score 17;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nero
                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PS,
                                                                                                                                                                                                                                                                                                                                                                   DB
69;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaarde
                                                                                                                                                                                                                                                                                                                                                                                     ۲,
                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                   Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               contacting cells with apoptotic disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WA;
                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                     Gaps
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sapiens.

Synthetic

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RESULT 43
AAX34894/c
ID AAX348
XX AAX348
XX AAX348
XX DE PCR pr
XX Immort
KW Immort
KW SV40 v
KW hair g
OS Synthe
XX Synthe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC can self-renew in the AFCI culture medium and can differentiate to CNS cereals self-renew in the AFCI culture medium and can differentiate to CNS cereals cells but not to CNS neuronal cells. The stem cells can be used to CC generate a population of mammalian motor neurons by incubating the stem cells in a medium promoting cell proliferation and neuronal cells in a medium comprises laminin-coated plates and NEP comedium lacking chick embryo extract. The stem cells can also produce neural crest stem cells by inducing the cells to differentiate in vitro. The inducing step comprises replating the cells to a laminin-coated comprises replating the cells on a laminin-coated comprise stem cells of the comprise comprise conding a dorsalizing agent to the cells, preferably fibroblast comprise conduce cells (BMP) such as BMP-2, -4 or -7. The stem cells can be used to produce cells of the peripheral nervous system, by inducing the stem cells to differentiate in vitro to neural crest stem cells, and inducing these cells to differentiate in vitro to neural crest stem cells, primers cells, and inducing these cells to differentiate in vitro to neural crest stem cells, and inducing comprise cells to differentiate in vitro to neural crest stem cells, and inducing comprise cells to differentiate in vitro to neural crest stem cells, and inducing comprise cells and chick embryons to have one cells and inducing comprise cells and chick embryons and chick embryons and inducing comprise cells and chick embryons and cells, and inducing comprise cells and chick embryons are cells and chick embryons and chick embryons and chick embryons and ch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                 Immortalized human hair papila cell; HPC; screening; hair growth; SV40 viral Large T-antigen gene; deleted replication initiation point; hair growth stimulating agent; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               differentiation to central nervous system (CNS) neuronal or glial cells and to neuronal crest stem cells. Also described is an isolated population of mammalian CNS glial-restricted precursor (GRP) cells which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes an isolated, pure population of mammalian neuroepithelial stem cells, which are capable of self-renewal in adherent feeder-cell-independent (AFCI) culture medium and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalian neuroepithelial stem cells and glial restricted precursor - can self renew and differentiate into central nervous system cells, used for generating various types of cells
                                                                                                                                                                    PCR primer used to
                                                                                                                                                                                                                    28-JUN-1999
                                                                                                                                                                                                                                                                                                                 AAX34894 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 26; Page 61; 78pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mayer-Proschel M, Mujtaba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-MAY-1998;
07-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1873 GAGATGGAGATGATGAA 1889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                        GAGATGGAGATGATGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP;
                                                               Large T-antigen gene; deleted restimulating agent; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        UTAH RES FOUND.
                                                                                                                                                                                                                 (first entry)
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97US-0852744
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; 1 C; 8 G; 5 T; 0 other;
                                                                                                                                                                 amplify FGFR3
                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
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                                                                                                                                                                                                                                                                                                                 20
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rao MS;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 22;
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RESULT 44
AAA95391/c
ID AAA95391 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The specification describes the preparation of immortalized human hair papila cells (HPC). The method comprises transformation of HPC with an SV40 viral Large T-antigen gene with deleted replication initiation point. The immortalized HPC can be used in a screening method for a hair growth agent, by culture of immortalized HPC in the presence of a substance to be tested and observation of the growth of the immortalized HPC. HPC is also used in development of hair growth stimulating agents. The present sequence represents a PCR primer, which is used in the course of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immortalized human hair papila cells used for evaluation growth agent - are prepared by transformation of human ha cells with gene with deleted replication initiation point
                        Cell comprising exogenous nucleic acid inducing tyrosine hydroxylase expression useful for treating catecholamine-related diseases such as Parkinson's disease, manic depression and schizophrenia -
                                                                                                                                                                                                                                                                                                    Rat; Nurr1; tyrosine hydroxylase; catecholamine-related disease; Parkinson's disease; manic depression; schizophrenia; PCR primer;
Example 1; Page 20; 68pp; English
                                                                                                                                                                                                                      05-OCT-2000.
                                                                                                                                                                                                                                                                          Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                            Rat FGFR coding sequence PCR primer #2.
                                                                                  WPI; 2000-656165/63
                                                                                                           Sakurada K,
                                                                                                                                      (SALK ) SALK INST BIOLOGICAL STUDIES.
                                                                                                                                                                 26-MAR-1999;
                                                                                                                                                                                           21-MAR-2000; 2000WO-US07544
                                                                                                                                                                                                                                                 WO200058451-A1.
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                                                                                                           Palmer T,
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hair papil
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes the rat Nurr1 coding and protein sequences. The Nurr1 protein is involved in the induction of tyrosine hydroxylase expression in adult rat-derived hippocampal progenitor cells. The Nurr1 gene and protein can be used in the treatment of catecholamine-related diseases such as Parkinson's disease, manic depression and schizophrenia. They can also be used to induce tyrosine hydroxylase expression and identify tyrosine hydroxylase related deficiencies, which are linked to the same diseases. The present sequence is a PCR primer used in a method to differentiate adult neural progenitor
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17; Conserv
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on; therapy; human; phosphorothioate; ss.
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Pred. No. 79
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01-MAR-2001; 2001US-0798096

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer; GPRalphal; GPI-anchored GDNF family receptor alpha-1; GDNF; glial cell line-derived neurotrophic factor; glycophosphatidyl-inositol; GPI; auditory neuron protector; improve learning ability; treatment; Parkinson's disease; epilepsy; neuronal disease; Ret; ss.
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                                                                                                                                                                                                                                                                                                                                                               01-OCT-1999;
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                             Methods for
                                                                                 WPI; 2000-317748/27
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screening for agonists and intracellular signalling pa
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Pred. No. 75;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
GPI-anchored alpha1-dependent,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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treatment of neuronal diseases
                                                                          Ret-independent intracellular signalling, useful in methods
                                                               71pp;
                                                                            for
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of receptor tyrosine kinase (Ret) expression. GDNP (glial cell line-derived neurotrophic factor) is a transforming growth factor-beta family related neurotrophic protein. GDNP action is thought to act through GBP-anchored GDNP family receptor alpha-1 (GPRalphal) and Ret. The PCR primer represented by this sequence is used in an example of the method of the invention. The method is used to identify compounds that are agonists or antagonists of intracellular signalling effected by glycophosphatidyl-inositol (GPI)-anchored receptors in cells of the nervous system. Ret-independent signalling gathways may stimulate neurotransmitter synthesis. The compounds identified by the method of the invention can be used for protecting auditory neurons, improving learning ability, treating epilepsy and treating Parkinson's disease. The present sequence represents a PCR primer used in the RT-PCR analysis tyrosine kinase (Ret) expression. GDNF (glial cell

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Query Match
Best Local
                              Matches
              2319
                              18;
20
                                     Similarity
TCAAAGTGATGTGTGGTCCT 1
             TCAGAGTGATGTCTGGTCCT 2338
                              Conservative
                                   1.2%;
                             0,
                                     Score 16.8;
Pred. No. 8
                              Mismatches
                                      90;
                                            BB
                                            1,
                             2
                                            Length
                              Indels
                                             21
                             0;
                              Gaps
                              0
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Sequence 21 BP; 7 A; 6 C; 4 G; 4 T; 0 other;

밁 S

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ARESULT 47
AAC92592/c
ID AAC925
XX AAC92592/c
XX AAC925
XX AAC925
XX AAC925
XX AAC925
XX Human
XX Human
XX Human
XX Human
XX Human
XX Human
XX Signal
XW Signal
XW Signal
XW System
XW Immune
XX Imm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cell growth; transcriptional repression; replication; signal transduction; chromatin decondensation; Ag-NOR fanucleolin antibody; systemic connective tissue disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human nucleolin; P92; C23; phosphoprotein; ribosome biogenesis; ribosome transport; cytokinesis; nucleogenesis; cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC92592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          scleroderma-like chronic graft versus host disease; expression inhibition; tumour formation; cancer; in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human nucleolin phosphorothioate antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              systemic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ag-NOR family;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SLE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IJ
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US6165786-A. Homo sapiens

mmune disorder;

phosphorothioate; antisense oligonucleotide; ss.

03-NOV-1999; 03-NOV-1999; 26-DEC-2000. 9908-0433699 9908-0433699

CF, SISI Cowsert LM; PHARM INC.

WPI; 2001-079848/09

hybridizes with for modulating t Claim 14; Column 41-42; 41pp; English Novel antisense compound targeted to human nucleolin which specifically the and inhibits the the expression of expression nucleolin i 'n of human nucleolin, useful in cells -

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AAZ59320/c
IID AAZ593
XX
AC AAZ593
XX
AC AAZ593
XX
AC Human
DE Human
XX
Single
KW Single
KW Single
KW Single
KW Single
KW Single
KW WOOD6;
XX
PT Variat
FT Variat
FT Variat
FT O9-JUN
XX
AC AXYS-
XX
PR (AXYS-
XX
PR 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            growth, transcriptional repression, replication, signal transduction, and chromatin decondensation. Nucleolin is a member of the Ag-NOR (active ribosomal gene located in the nucleolar organiser region) family of proteins which are markers of active ribosomal genes, and whose expression is associated with the prediction of tumour growth rate. The presence of antibodies against nucleolin are associated with systemic connective tissue diseases such as systemic lupus erythematosus (SLE) and scleroderma-like chronic graft versus host disease. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequences AAC92560-C92639 represent antisense oligonucleotides targetted to the human nucleolin gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human nucleolin mRNA, and were analysed for their effect on nucleolin mRNA levels by quantitative real-time PCR. Nucleolin (also known as P92 or C23) is the most abundant nucleolar phosphoprotein in actively growing cells. Nucleolin primarily participates in ribosome biogenesis and transport of ribosome to provide the primarily participates in ribosome biogenesis and transport of ribosome to provide the primarily participates and transport of ribosome being able to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transiently bind to pre-ribosomes in the nucleolus via a ribonucleoprotein consensus sequence. However, it has also been shown to be involved in cytokinesis, nucleogenesis, cell proliferation and
                                    Claim 2; Page 16; 46pp; English
                                                                                                              Novel nucleic acid used for genotyping, metabolism -
                                                                                                                                                                                                                      WPI; 2000-105892/09
                                                                                                                                                                                                                                                                                    Guida M,
                                                                                                                                                                                                                                                                                                                                                                                                                                    10-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-DEC-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9964630-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Single nucleotide polymorphism; SNP; STP2; phenol sulphotransferase; probe; genotyping; human; drug metabolism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human STP2 exon 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ59328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ59328 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligonucleotides of the invention are useful for diagnosis, prevention and treatment of conditions associated with nucleolin expression, such
                                                                                                                                                                                                                                                                                                                                                                  (-SYXA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1878 GGAGATGATGAAGATGAT 1895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19
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17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                formation,
                                                                                                                                                                                                                                                                                                                                                                  AXYS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GAAGATGATGAAGATGAT
                                                                                                                                                                                                                                                                                    Kurth J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                  PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0088710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99WO-US13094.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag=
/note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymorphism sequence 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immune disorders and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a
"Site of polymorphism'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 16.4; D
Pred. No. 86;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T; 0
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                                                                                                                                            e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB
                                                                                                                                               to predict rate of drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                      This invention describes novel Staphylococcus-specific oligonucleotides based on the consensus femA nucleotide sequence which are used to develop products for the identification, detection and therapy of infections. The oligonucleotides can be used for the genetic amplification, the identification and/or quantification of various femA sequences which are specific to known or unknown Staphylococci species.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acetaminophen, and paranitrophenol. Several of the nucleotide changes identified at the polymorphism sites, give rise to an amino acid change. Amino acid changes to the polymorphism sites, give rise to an amino acid change. Amino acid changes may result in altered enzyme activity. The sequences can be used as probes for detecting STP2 polymorphisms. The polymorphic probes are used in screening and genotyping, i.e. to predict the rate of metabolism of STP2 substrates, potential drug-drug interactions and adverse side effects. They can also be used to detect diseases resulting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequences AAZ59305-Z59352 are fragments of the human STP2 gene. fragments are from the 8 exons, the promoter region, 3' and 5' untranslated regions of the STP2 gene Bach sequence contains a identified STP2 gene single nucleotide polymorphism (SNP). STP2 phenol sulphotransferase. Substrates for STP2 include minoxidil, phenol sulphotransferase. Substrates for STP2 include minoxidil,
              Since the femA sequence is similar to the femB sequence, the oligonucleotides can also be used for the molecular genotyping of femB genes of different Staphylococci species or other gram-positive bacter. The femA nucleic acids can also be used in therapeutic applications. They can also be used to identify inhibitors, e.g. antibodies or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX37791 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 8 A; 6 C; 4 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FemA; primer; i
amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX37791;
                                                                                                                                                                                                                                   Example 1; Fig 1; 48pp; English.
                                                                                                                                                                                                                                                                     New Staphylococcus-specific oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                  26-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Staphylococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                               (AXTO-)
                                                                                                                                                                                                                                                                                                                                                                                                (BENA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    accidental or occupational exposure to toxins and to establish al, cell or in vitro models for drug metabolism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18
                                                                                                                                                                                                                                                                                                                                                                             BELGIAN MIN NAT
UNIV CATHOLIQUE
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                                                                                                                                                                                                                                                                                                                                              Vannuffel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CATAGGGGTCTTCTTCAT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     identification; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                genotyping; gram-positive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98WO-BE00141.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94.4%;
                                                                                                                                                                                                                                                                                                                                                                                                DEFENCE
                                                                                                                                                                                                                                                                                                                                                                               LOUVAIN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        oligonucleotide
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Pred. No. 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  therapy; infection; bacteria; vaccine; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
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for blocking expression

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                                               The sequences given in AAQ51825-2266 represent areas of mRNAs which associated with development or maintenance of chronic myelogenous leukemia (CML), promyelocytic leukemia, Burkitt's lymphoma, or acute lymphocytic leukemia, follicular lymphoma, B-cell acute lymphocytic leukemia, breast cancer, colon carcinoma, neuroblastoma and lung cancer. The full length mRNAa containing themes to contain the containing themes the containing the containing themes the containing the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Multiple drug resistance; mdr-1; ribozyme; membrane protein; liver; resistance; chemotherapeutic agent; colchicine; doxorubicin; colon; actinomycin D; vinblastine; small intestine; kidney; adrenal gland; adenocarcinoma; bowel; transformed phenotype; promyelocytic leukemia; human; chronic myelogenous leukemia; CML; follicular lymphoma; B-cell acute lymphocytic leukemia; breast cancer; colon carcinoma, neuroblastoma; lung cancer; genetic drift; mutation; hammerhead motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleotide sequences.
against Staphylococci
    sequences, encode aber
cellular proliferation
                                                                                                                                                                                                          Claim 3; Fig 10; 69pp; English.
                                                                                                                                                                                                                                                       multiple drug resistance
                                                                                                                                                                                                                                                                            New enzymatic RNA molecules (ribozymes) - associated with tumours or mRNA expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-AUG-1992
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26-AUG-1992;
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14-MAY-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hairpin; hepatitis
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26-MAY-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                            RIBOZYME PHARM INC
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93US-0006122.
93US-0008910.
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92US-0936531.
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92US-0882885.
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The full length mRNAs containing these target aberant cellular proteins which are able to cation and are directly linked to a leukemic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.2%;
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Pred. No. 99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB
                                                                                                                                                                                                                                                                       from gene encoding
                                                                                                                                                                                                                                                                                                    which cleave mRNA
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Query Match Best Local Similarity

Score 15.8; DB 1; Pred. No. 1e+02;

Length

19;

19

BP;

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5 C; 1.1**%**; 89.5**%**;

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other;

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restenosis treatment.

designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDKI, PCNA and Cyclin BI. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787 The ribozyme of the invention is useful for

The present invention relates to a hairpin or hammerhead ribozyme,

AAA8415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is

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RESULT 51
AAA82642
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   resistance (mdr-1) mRNA specific ribozymes remove the mechanism of drug resistance used by transformed cells and thus enhances drug therapies for tumours. The ribozymes may also be used to study genetic drift and mutations within cells.

(Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the invention. The ribozymes is formed in a hammerhead motif, but may also be formed in the motif of a hairpin, hepatitis delta virus, group I intron or RNaseP-like RNA. These ribozymes may be used to inhibit the development or expression of a transformed phenotype in man and other animals by modulating expressed in pre-neoplastic and transformed cleavage of target mRNAs expressed in pre-neoplastic and transformed cells elicits inhibition of the transformed state. Multiple drug
                                                                                                                               Disclosure; Page 49; 109pp; English
                                                                                                                                                     New hairpin and hammerhead ribozyme RNA encoding a cyclin or cell-cycle PCNA and Cyclin B1 -
                                                                                                                                                                                                        WPI; 2000-412314/35
                                                                                                                                                                                                                                                                              04-DEC-1998;
                                                                                                                                                                                                                                                                                                      06-DEC-1999;
                                                                                                                                                                                                                                                                                                                              08-JUN-2000
                                                                                                                                                                                                                                                                                                                                                     WO200032765-A2
                                                                                                                                                                                                                                                                                                                                                                              Mammalia
                                                                                                                                                                                                                                                                                                                                                                                                                Ribozyme; hairpin; hammerhead; gene therapy; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                      cdk2 ribozyme binding site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 8
                                                                                                                                                                                                                              Tritz R,
                                                                                                                                                                                                                                                       (IMMU-) IMMUSOL INC
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                                                                                                                                                                                                                              Welch PJ,
                                                                                                                                                                                                                                                                                                                                                                                                      88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                              Barber JR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 16;
Pred. No.
                                                                                                                                                                                                                              Robbins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 other;
                                                                                                                                                                   for inhibiting restenosis, cleaves dependent kinase other than CDK1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 19;
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RESULT 52
AAH57804
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                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, antisickling dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; head cell carcinoma;
                                                                  also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of grematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scarr. AhH57577 to AhH62099 represent sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent
                                                                                                                                                                                                                                            dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can appear to the contraction of the contraction
                                                        exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 88; 408pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-300427/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cell-cycle dependent kinase cdk2 ribozyme binding site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTTTGGACTAGCCAGAGCT 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tritz R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0161532
                                                        present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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Sequence 19

BP; 4

A; 5 ç;

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υī

T; 0 other

HERG gene exon 7/intron 7 junction sequence

19-JUN-2000

(first entry)

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RESULT 54
AAA07637/c
ID AAA076
XX
AC AAA076
XX
DT 19-JUN
XX
DE HERG G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 53
AAZ07940/c
ID AAZ079
XX
AC AAZ079
XX
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                                                                                                                                                                        Matches
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                              useful for the treatment of asthma and in gene the ripper. IZEC-CIM is of allergic diseases. IZEC-CIM blocks IgE binding to mast cells/basophils 10000 fold better than native soluble CD23, and has comparable blocking and binding efficiency to antibodies, but without any side effects. Sequences AAZ07939-946 represent PCR primers used in the preparation of chimeric CD23 constructs.
                                             AAA07637;
                                                                  AAA07637
                                                                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ07940 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                             Disclosure; Column 4; 18pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-579942/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chimeric protein; IzEC-C1M; extracellular; CD23; asthma; gene therapy; allergic disease; mannose binding protein; MBP; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mannose binding protein (MBP) cDNA fragment amplifying primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ07940;
                                                                                                                                                                                                                                                                                                                                                                              Chimeric protein derived from the extracellular region of CD23, useful
                                                                                                                                                                                                                                                                                                                                                                                                                            Conrad DH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US5965712-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                   for treating asthma and allergic diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYVI-) UNIV VIRGINIA COMMONWEALTH
                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2205
                                                                                                                                        1819 GTGAAGATGTTGAAAGATG 1837
                                                                                                                                                                       l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 CTTTGGACTAGCCAGAGCT 19
                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTTTGGACTCGCCAGAGAT
                                                                                                                           GTGAAAATGTTGAAAGAAG
                                                                                                                                                                                                                    BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                           Kelly AE;
                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0100398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98US-0100398
                                                                   DNA;
                                                                                                                                                                                 89.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.1%;
                                                                                                                                                                                                                    6 C; 0 G; 10 T; 0 other;
                                                                   20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2223
                                                                                                                                                                                             Score 15.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.8; DB 1;
Pred. No. 1e+02;
                                                                                                                                                                                    Pred.
                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                   NO.
                                                                                                                                                                                 1.1e+02;
                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2
                                                                                                                                                                                            Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                        Indels
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                                                                                                                                                                        Gaps
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a HERG protein having a mutation compared to wild-type HERG, and is useful for developing products for the diagnosis, prevention and treatment of long QT (LQT) syndrome. The products and methods can be used for the diagnosis of subjects with LQT syndrome. They can also be used to screen for drugs for treating or preventing LQT syndrome. The HERG nucleic acids can also be used for gene therapy and HERG peptides can be used for peptides can be used for gene therapy.
                    16-MAY-2000;
07-DEC-2000;
                                                                                                                                                                       Telomerase reverse transcriptase; TERT; cytostatic; cell growth inhibitor; antisense oligonucleotide;
                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                   09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-JUL-1998;
06-JAN-1999;
(ISIS-) ISIS PHARM INC
                                                       15-MAY-2001; 2001WO-US15774.
                                                                                22-NOV-2001.
                                                                                                      WO200188198-A1
                                                                                                                                                                                                           Telomerase reverse transcriptase, antisense oligonucleotide
                                                                                                                                                                                                                                                          AAS96638
                                                                                                                                                                                                                                                                                 AAS96638 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 2 A; 4 C; 7 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 8; Page 71; 163pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-195319/17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Keating MT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UTAH ) UNIV UTAH RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HERG; mutation; long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        isolated mutant HERG nucleic acids, useful the diagnosis, prevention and treatment of
                                                                                                                                                                                                                                                                                                                                                               2556 CACTCTCACAACCAATGAG 2574
                                                                                                                                                                                                                                                                                                                                          19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88.
                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis, prevention
                                                                                                                                                                                                                                                                                                                                          CACACTCACAGCCAATGAG 1
                                                                                                                                                              technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          intron/exon junction sequences of the
                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                       2000US-0733294.
                                  2000US-0572423.
                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Splawski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0122847.
99US-0226012.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US16337.
                                                                                                                                                                                                                                                                                                                                                                                                 1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        QT syndrome; LQT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ţ,
                                                                                                                                                                                                                                                                                 ВÞ
                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                             Score 15.8; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                    Pred.
                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                    1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                      2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for developing products long QT syndrome -
                                                                                                                                                                                                                                                                                                                                                                                                                                                            HERG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                           Length 20
                                                                                                                                                                                  apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene.
                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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WPI; 2002-075321/10. Monia BP, Gaarde WA, Freier , MS Wancewicz

expression of TERT, useful for modulating New compound targeted to nucleic acid molecule encoding telomerase transcriptase (TERT), which specifically hybridises with and inhibits apoptosis and inhibiting cell

Claim 26; Page 91; 154pp; English.

CC transcriptase, where the compound specifically hybridises with and CC inhibits the expression of TERT. A series of oligonucleotides were CC designed to target different regions of the human TERT RNA. These were CC 20 nucleotides in length and composed of a central gap region consisting CC of ten 2'-deoxynucleotides, flanked on both sides (5' and 3' directions) CC by five-nucleotide wings. The wings were composed of 2'-methoxyethyl CC (2'-mOE) nucleotides. The compounds were analysed for their effect on human TERT mRNA levels by reverse transcriptase (RT)-polymerase chain CC reaction (PCR). The compound is useful for inhibiting the expression of CC TERT in cells or tissues, for treating a human having disease or CC condition associated with TERT, for modulating apoptosis, for inhibiting CC cell growth (preferably, cancer cell growth), in antisense therapy and CC coligonucleotide used to modulate the activity of nucleic acid molecules CC encoding TERT, described in the method of the invention. to a nucleic acid molecule encoding human TERT (telomerase reverse The invention describes a compound, 8-50 nucleobases in length targeted

Sequence 20 BP; 4 A; 3 C; 6 G; 7 T; 0 other;

Ş Matches Query Match Best Local Local 2020 GGGATGGAGTACTCCTATG 2038 l Similarity 17; Conserv Conservative 1.1%; 0 Score 15.8; Pred. No. 1. Mismatches 1.1e+02; hes 2; DB 1; Length 20; Indels 0 Gaps 0

밁 GGGATGGACTATTCCTATG 19

RESULT 56

ABS97467 standard; DNA; 21

23-DEC-2002 (first entry)

Human diazepam binding inhibitor (DBI) gene polymorphic sequence #11.

ABS97467/c
ID ABS974
XX
AC ABS974
AC ABS974
XX
XX
AC ABS974
XX
XX
BY
C 23-DEC
XX
C Human
XX
KW Human;
KW Cycloo
KW adrene
KW aryl h
KW cycloo
KW epoxid
KW glutat
KW HNWT;
KW HNWT;
KW WADPH
KW WOTH
KW WATH
K KW Human; ds; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;
KW cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF;
KW adrenergic receptor beta1; ADBR1; arryl hydrocarbon; AHR; MRP3; NR112;
KW aryl hydrocarbon receptor nuclear translocator; ARNT; Cathepsin S; CTSS;
KW cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;
kW epoxide hydroxylase 2; EPHX2; 5-11poxygenase activating protein; FLAP;
KW glutathione-S-transferase 12; GST12; histamine-N-methyl transferase;
KW HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase;
KW MADPH quinone oxidoreductase 2; NOO2; sulfotransferase thermolabile;
KW MADPH quinone oxidoreductase 2; NOO2; sulfotransferase thermolabile;
KW MADPH quinone oxidoreductase 2; NOO2; sulfotransferase receptor; uPA;
KW MUT; UDP-glucuronosyl transferase 2B7;
W UGT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;
KW multidrug resistance 1; lactotransferain; orphan nuclear receptor; uPA;
KW multidrug resistance associated protein 3; cancer; prostate;
KW multidrug resistance associated protein 3; cancer; prostate;
KW acetylcholine muscarinic receptor; UGMRA; CHMR3; CHMR4; CHMR5;
KW altered drug metabolism; cardiovascular function; colorectal tumour;
KW central nervous system; pulmonary; immunological; SNP; polymorphism

WO200257410-A2

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CC cyrochrome P430 AI (CYP450LAI), Cyrochrome P430 AZ (CYP450CEI), adrenergic receptor betal (ABRI), CC aryl hydrocarbon (AFR), aryl hydrocarbon receptor nuclear translocator (C (ARNT), cathegisin S (CTSS), cyclooxgenase 2 (COX2), diazepam binding C inhibitor (DBI), epoxide hydroxylase 2 (EPHX2), 5-lipoxygenase C inhibitor (DBI), epoxide hydroxylase 2 (EPHX2), 5-lipoxygenase C inhibitor (DBI), epoxide hydroxylase 2 (EPHX2), 5-lipoxygenase C inhibitor (DBI), protein (FLAP), glutathione-S-transferase 12 (GST12), the transferase (INMT), cathegial (INTR), cathegial (INTR), sellikrein 2) KLZ2, nicothiamide C insentine-N-methyl transferase (INMT), (kallikrein 2) KLZ2, nicothiamide C Insentine-N-methyl transferase (INMT), Label (INTR), multidrug resistance 284 (UGT1284), UDP-glucuronosyl transferase 284 (UGT1284), UDP-glucuronosyl transferase 284 (UGT1287), UDP-glucuronosyl transferase 287 (UGT1287), UDP-glucuronosyl transferase 284 (UDP-glucuronosyl tran
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                                   á
                                                                                    Matches
                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related traits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-698522/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-NOV-2001; 2001WO-US44838
                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to the sequence of an isolated nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 9; Page 115; 714pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (DNAS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-NOV-2000; 2000US-0724389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-JUL-2002.
                                                                                                                                                                                                                sequence
                                          2055 TGAGGAGCAGATGACCTTC
                                                                                    17;
                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA SCI LAB INC
                                                                                                                                                                        21
TGATGAGCAGATGCCCTTC 2
                                                                                                                                                                        BP; 6
                                                                                    Conservative
                                                                                                                                                                                                                  of the
                                                                                                                                                                     A; 5 C; 6
                                                                                                                                                                                                                  invention.
                                                                                                   1.1%;
                                                                                    <u>.</u>
                                                                                                                                                                        G; 4 T; 0
                                                                                                     Score 15.8; DB 1;
Pred. No. 1.1e+02;
                                                                                    Mismatches
                                                                                                                                                                        other;
                                                                                    1.1e+02;
ches 2;
                                                                                                                         Length 21;
                                                                                    Indels
                                                                                    0;
                                                                                    Gaps
                                                                                    0
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KW cytochrome P450 A2; CYP4501A2; Cytochrome P450 02E; CYP45002E1; LTF; KW adrenergic receptor beta1; ADBR1; aryl hydrocarbon; AHR; MRP3; NR112; KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS; KW cycloxygenase 2; CCX2; diazepam binding inhibitor; DBI; haematological; KW cycloxygenase 2; CCX2; diazepam binding inhibitor; DBI; haematological; KW epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP; KW glutathione-S-transferase 12; GST12; histamine-N-methyl transferase; MNMT; Kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT; KW MADPH quinone oxidoreductase 2; NQO2; sulfotransferase thermolabile; STM; UDP-glucuronosyl transferase 2B4; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA; W STM; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA; W multidrug resistance 1; lactotransferance; prostate; RW multidrug resistance associated protein 3; cancer; prostate; RW multidrug resistance associated protein 3; cancer; prostate; RW acetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5; RW altered drug metabolism; cardiovascular function; colorectal tumour; RW central nervous system; pulmonary; immunological; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human epoxide hydroylase 2 polymorphic sequence #46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ds; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;
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25-JUL-2002.

28-NOV-2001; 2001WO-US44838.

28-NOV-2000; 2000US-0724389.

(DNAS-) DNA SCI LAB INC

Guida M, Hall J;

WPI; 2002-698522/75.

Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related traits genetic linkage the genes

Example 10; Page 118; 714pp; English.

CC This invention relates to the sequence of an isolated nucleic acid CC molecule comprising at least one base variation from that of a known CC human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A2), CC cytochrome P450 02E1 (CYP4502E1), advenergic receptor beta1 (ADBR1), CC aryl hydrocarbon (AHR), aryl hydrocarbon receptor nuclear translocator (C ARNT), cathepsin S (CTGS), cyclooxgenase 2 (COX2), diazepam binding cinhibitor (DBI), epoxide hydrocylase 2 (EPHX2), 5-lipoxygenase cc inhibitor (DBI), epoxide hydrocylase 2 (EPHX2), 5-lipoxygenase cc inhibitor (DBI), epoxide hydrocylase 2 (EPHX2), 5-lipoxygenase cc activating protein (FLAP), glutathione-S-transferase 12 (GST12), cc lipoxygenase (NDMT), (Kallikrein 2) KLK2, nicotinamide cc lipoxygenase cc (NDMT), NADPH quinone oxidoreductase 2 (NQO2), cc sulfotransferase thermolabile (STM), UDP-glucuronosyl transferase 2B4 (UGT2B4), UDP-glucuronosyl transferase 2B7 (UGT2B7), UDP-glucuronosyl transferase receptor (NPA), multidrug resistance cc (MT2B1), orphan nuclear receptor (NPA), or acetylcholine cc (MT2B1), orphan nuclear receptor (NRI12), or acetylcholine cc (MRS1) sequence. The polymorphisms in the human genes cited in the convertion are useful as genetic linkage markers for locating and cc the genome and eventually identifying the genes responsible for a cc variety of disorder-related traite as a result of their e-g., contained covertopression, constitutive expression, mutation or underexpression, which may be used in diagnosing and/or treating the disorders. The nucleic acid molecules comprising the polymorphic sequences contained in CYP4501A1, CYPP4501A2, CYP4502E1, ARNT, EPHX2, GST12, NNMT, NQO2, NR112, STM, UGT2B4, UGT2B7, UGT2B15, AHR, MDR1 and/or MDR3 are useful

ABS97555 ID ABS9

ABS97555 standard; DNA; 21 BP

RESULT 57

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RASULT 58
ABQ84277
ID ABQ84
AC ABCA
AC A
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Best Local :
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24-APR-2001;
12-OCT-2001;
12-OCT-2001;
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                                                                                  The present invention describes a new isolated nucleic acid sequence (I) comprising a DPP10 mRNA sequence. DPP10 is a dipeptidyl peptidase (also known as prolyloligopeptidase). (I) has antiinflammatory, antiasthmatic,
                                                                                                                                                                                                                                                                                                     New nucleic acid sequence comprising DPP10 mRNA, useful for the manufacture of a medicament for regulating DPP10 protein expression for preventing or treating inflammatory disease e.g., inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DPP10; dipeptidyl peptidase; prolyloligopeptidase; enzyme; asthma; antiinflammatory; antiasthmatic; antipsoriatic; antiarthritic; antitheumatic; vaccine; gene therapy; inflammatory disease; inflammatory bowel disease; atopy; rheumatoid arthritis; psoriasis
                                                                                                                                                                                                                     Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-APR-2002;
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                                                                                                                                                                                                                                                                                 preventing
el disease
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; 2001GB-0010046.
; 2001GB-0024575.
; 2001GB-0024594.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INNOVATIONS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Moffat MF, Allen
                                                                                                                                                                                                                     70;
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                                                                                                                                                                                                               321pp;
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                                                                                                                                                                                                           English
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Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Σ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rheumatoid arthritis; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lench
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antipeoriatic, antiarthritic and antirheumatic activities, and can be used in vaccines and gene therapy. A composition comprising (I) can be

in vaccines and gene therapy. A composition comprising (I) can be for the manufacture of a medicament for regulating DPP10 expression

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RESULT 59
AAX74847
ID AAX7
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or for preventing or treating inflammatory disease e.g., inflammatory bowel disease, asthma, atopy, rheumatoid arthritis or psoriasis. (I) c also be used in an assay for detecting or measuring DPP10 in a sample. A host cell comprising (I) can be used for producing recombinant DPP10 gene products, or in drug screening systems to identify agents for diagnosis or treatment of individuals having or susceptible to inflammatory disease. Human DPP10 is located on chromosome 2, more specifically chromosome 2q14. ABQB4254 to ABQB4612 and ABP55569 to ABP55629 represent sequences used in the exemplification of the present invention.
          The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. ANX67275 to ANX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                     Nucleic acid molecule modulating VEGF receptor(s) mRNA stability - useful for treating e.g. tumour psoriasis, rheumatoid arthritis, etc., in a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage;
tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease;
fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                      Claim 4; Page 166; 218pp; English. .
                                                                                                                                                                                                                                                             WPI; 1997-259017/23.
                                                                                                                                                                                                                                                                                        Escobedo J,
                                                                                                                                                                                                                                                                                                                                                               11-JAN-1996;
26-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                           25-OCT-1996;
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flk-1; KDR;
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17; Conserv
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                                                                                                                                                                                                                                                                                                                      RIBOZYME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard; RNA;
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95US-0005974.
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                                                                                                                                                                                                                                                                                                                      PHARM INC.
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Pred. No. 1.1e
0; Mismatches
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RESULT 60
ANY96481/c
ID AAV964
XX AAV964
XX AV964
XX DO 1-MAR
XX Solant
KW Hammer
CO Solant
KW 24-NOV
PR 28-JAN
XX 24-NOV
PR 28-JAN
XX Claim
XX New en
PT McSwig
DR WPI; 1
XX New en
PT Biosyr
XX Claim
XX Clai
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                                                       alkaloids; or (ii) involved in flower formation. AAV9592 to AAV96334, cand AAV96335 to AAV96354 represent potate solanidine glucosyltransferase hammerhead and hairpin ribozymes, respectively. AAV95629 to AAV95981, and AAV96355 to AAV96734 represent potato solanidine glucosyltransferase target sequences. AAV96773 to AAV97170, and AAV97171 to AAV97195 represent potato citrate synthase hammerhead and hairpin ribozymes, respectively. AAV96773 to AAV97772, and AAV97196 to AAV97120 represent potato citrate synthase target sequences. Ribozymes of the present invention can be used to inhibit the synthests of toxic alkaloids in solanaceous plants, particularly potato but also tomato, pepper, aubergine and ditura or to inhibit flowering in potato, lettuce, spinach, cabbage, brussel sprouts, arugula, kale, collards, chard, beet, turnip, sweet potato and turf grass. Also the ribozymes can be used for RNA manipulation in the same way that restriction endonucleases are for DNA, as well as to examine genetic drift and mutations in plants and to consensus sequences within a family of related genes, and being catalytic need to be present at only very low concentrations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-NOV-1997;
28-JAN-1997;
28-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Solanidine; glucosyltransferase; potato; citrate synthase; target; hammerhead ribozyme; hairpin ribozyme; alkaloid biosynthesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNA-cleaving activity (e.g. ribozymes) which are capable of modulating the expression of plant genes: (i) involved in biosynthesis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes enzymatic nucleic acid molecules with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 53; Page 53; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New enzymatic nucleic acid(s) - useful for, e.g. reducing alkaloid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                McSwiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9832843-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Solanum tuberosum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Potato citrate synthase target sequence position 543
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV96481;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV96481 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GAAGAUGUUGAAAGAGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97US-0979416.
97US-0036545.
97US-0036599.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zwick MG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-US00738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; 0 C; 7 G; 3 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cleavage; solanaceous plant; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>ب</u>
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Pred. No. 99;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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Query Match

Sequence

17

BP; 4 A; 5 C; 2

G; 6 U; 0

other;

1.1%;

Score 15.4; DB 1;

Length 17

RESULT 62 AAX60990/c ID AAX609

AAX60990 standard; cDNA to mRNA; 19

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RESULT 61
ABQ81992/c
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                                                        Query Match
Best Local Similarity
Matches 16; Conserv
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                                                                                                                                                             The present invention describes a method for determining if an individual has a tumour cell or site of angiogenesis, or if a treatment is effective in changing angiogenesis or changing a status of a set of target cells, comprising angiogenesis or changing a status of a set of target cells, comprising determining if a sample of the subject has an expression product of at least one marker gene. Also described is a compound capable of altering the expression or activity of Keratin 14, TIE 1, Salioadhesin or Siglec in a cell. Peripheral blood mononuclear cell (PBMC)-expressed Keratin 14, TIE 1, Salioadhesin or Siglec, and Kits containing them from the present invention can be used in a diagnostic method, particularly as an indicator of angiogenesis or to determine presence of a tumour cell. The method of the invention is suitable to determine within a few days if a certain treatment against Kaposi's Sarcoma is successful. ABQ81851 to ABQ82006 represent nucleotide sequence used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                 Determining presence of a tumor cell or angiogenesis, and the effectiveness of treatment, by detecting the presence of marker is useful to detect and monitor treatment of Kaposi's Sarcoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABQ81992 standard;
                                                                                                                      Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; Kaposi's sarcoma; tumour; angiogenesis; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                      Example 10; Page 24; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-668396/72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-JAN-2001;
28-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-JAN-2002; 2002EP-0075264.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kaposi's Sarcoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (AMST-) AMSTERDAM SUPPORT DIAGNOSTICS
                            2413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2470 TACATGATGATGAGGGA 2486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17
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                                                                                                                                                     invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kuyl AC,
                    AAGCTGCTGAAGGAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TACATGATGATCAGGGA 1
                                                      1.1%;
nilarity 94.1%;
Conservative
                                                                                                                      B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; 2001EP-0200228.
; 2001EP-0203703.
; 2001US-325722P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                      2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TAG PCR primer SEQ ID NO:142.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cornelissen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA; 18
                                                                                                                      8 C; 3 G; 5 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>,</u>
                                                                         Score 15.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. 99;
                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                           .1e+02;
                                                                                        DB 1;
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                                                                                      Length 18;
                                                              Indels
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RESULT 63
ABK71996/c
ID ABK719
XX
AC ABK719
XX
AC ABK719
XX
DT 30-JUL
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                                                                                                                                                                                                                                                                                                                                                               CC result in a light hypersensitive phenotype. The gene, wen altered, is C responsible for the light hypersensitive mutant phenotype in Solanum CC lycopersicum (tomato) plants, the phenotype comprising a reduced growth CC of the plant associated with high levels of carotenoids and/or CC chlorophylls and/or flavonoids. Vectors comprising the sequence gene can be used to produce transgenic plants, such as pepper, egyplant, soybean, CC grape, melon, rice, carrot, spinach, citrus, pomaceae or ornamental comprised that contain a tomato TDETI gene. The TDETI mutants are useful con the agro-industrial sector, for generating tomato fruits with high CC carotenoid and/or flavonoid contents. Carotenoids and flavonoids have cantioxidant properties and in addition some flavonoids exhibit cantitumoural properties. They also exhibit a role in plant protection cagainst pathogenic agents and UV light irradiation. Menipulation of the TDETI gene expression can also be used to modify anthocyanin and CC carotenoid content in ornamental species for the achievement of new CC colour variants. Alteration of carotenoid content is useful for improving resistance to Norflurazon herbicides. Further it is possible to combine a content of feath view with mutations such as rin, nor and Nr, which
                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                           Query Match
Best Local
 30-JUL-2002
                                      ABK71996
                                                                       ABK71996 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleotide sequences of the tomato TDET1 (HP-2) gene, which modified, results in a light hypersensitive phenotype
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tomato; TDET1 (HP-2) gene; light hypersensitive phenotype; mutation; carotenoid; chlorophyll; flavonoid; transgenic; TDETI gene; anthocyanin; agro-industrial; antioxidant; antitumoural; plant protection; variant; ornamental; herbicide; fruit ripening; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                 Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention describes a tomato TDET1 (HP-2) gene, which if modified result in a light hypersensitive phenotype. The gene, wen altered, is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 15; 57pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-385610/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tomato TDET1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lycopersicon sp.
                                                                                                                                                                                                      1816 GCCGTGAAGATGTTGAA 1832
                                                                                                                                                                   19
                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                   GCCGTGAAGATGATGAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mustilli
                                                                                                                                                                                                                                                                                                                 BP; 4 A; 6 C; 3 G; 6 T; 0 other;
                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene amplifying nested primer.
 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97IT-RM00760
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                                                                                                                                                                                                                                                                                                                                                  ripening process.
                                                                                                                                                                                                                                                             94.1%;
                                                                                                                                                                                                                                          <u>.</u>.
                                                                                                                                                                                                                                                          Score 15.4; DB 1;
Pred. No. 1.2e+02;
                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                             Length 19;
                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                        0,
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cancer; DNA methylation; adenocarcinoma; leukaemia; lymphoma; melanoma; myeloma; sarcoma; teratocarcinoma; breast; prostate; liver; ovary; head and neck cancer; neuroectoderm; placenta; skeletal muscle; tonsil; lymph tissue; kidney; colon; uterus; testis; stomach; adrenal gland; bladder; bone; bone marrow; gall bladder; ganglia; salivary gland; gastrointestinal tract; parathyroid; penis; salivary gland; spleen; synovial membrane; thymus; thyroid gland; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                   Human MTG16 gene,
                                                                                                                                                                                                                                                                                                                           tumour suppressor gene;
                                                                                                                                                                                                                                                                                                                                                                                                                       PCR
                                                                                                                                                                                                                                                                                                                                                                                                                       primer #31.
                                                                                                                                                                                                                                                                                                                           5-aza-2'-deoxycytidine;
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Homo sapiens.

WO200218592-A1

31-AUG-2001; 2001WO-AU01097.

01-SEP-2000; 2000AU-0009806

(BION-) BIONOMICS LTD

Callen DF, Whitmore SA, Kremmidiotis G, Kochetkova M, Crawford ŗ,

Tumour suppressor gene which encodes polypeptide MTG16 active in suppressing cellular function associated with cancer, useful for manufacturing a medicament for treating cancer e.g. adenocarcino adenocarcinoma,

8; Page 67; 125pp; English.

The invention relates to an novel tumour suppressor gene (I) MTG16a or CC MTG16b and encoded polypeptide (II). (I) and (II), or a compound that CC mimics MTG16 activity such as 5-aza-2'-deoxycytidine, are useful for CC manufacturing a medicament for treating cancer, where it reverses DNA CC myeloma, sarcoma, teratocarcinoma, leukaemia, lymphoma, melanoma, CC myeloma, sarcoma, teratocarcinoma, and especially cancer of the breast, CC prostate, liver, ovary, head and neck, neuroectoderm, placenta, skeletal CC muscle, tonsil, lymph tissue, kidney, colon, uterus, testis, stomach, CC adrenal gland, bladder, bone, bone marrow, gall bladder, ganglia, CC gastrointestinal tract, parathyroid, penis, salivary glands, skin, spleen CC synovial membrane, thymus and thyroid gland. (I) and (II) are useful for CC diagnosis of a cancer, or predisposition to cancer, by establishing a CC primers derived from (I). MTG16 is further useful for identifying CC interacting protein suitable as drug targets, where MTG16 is fused to a CC DNA binding domain and used as the bait in a yeast two-hybrid system. CC ABK71964-ABK72041 represent human MTG16 coding sequences and PCR cc primers of the invention.

Sequence 19 BP; 2 A; 6 C; 6 G; 5 T; 0 other;

밁 Matches Query Match Local Similarity hes 16; Conserv 1935 CACACAGGATGGGCCTC 1951 17 CACACAGGATGGGCCAC Conservative 1.1%; 0 Score 15.4; DB 1; Pred. No. 1.2e+02; Mismatches ..2e+02; Length 19; Indels 0 Gaps

25-SEP-1996 AAT31581 standard; DNA; 20 BP (first entry)

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RESULT 65
AA169511/c
ID AA1695
XX AA1695
XX Intest
XX In
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Preventing or promoting programmed cell death in vertebrate cells comprises inhibiting or increasing the activity of interleukin-1-beta converting enzyme, or altering expression of
        19-OCT-1999;
                                                                                    19-OCT-1999;
                                                                                                                                                                 24-APR-2001
                                                                                                                                                                                                                                              JP2001112485-A
                                                                                                                                                                                                                                                                                                                           Bacteria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Intestinal bacteria detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAI69511;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAI69511 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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interleukin-1 bet
                                                                                                                                                                                                                                                                                                                                                                                                    Intestinal
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beta converting enzyme; gene therapy; primer; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95US-0368704.
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        99JP-0296815
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                                                                                                                                                                                                                                                                                                                                                                                                detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
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Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                    PCR
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RESULT 66
AAH20838
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Best Local
This invention describes a novel method for the direct identification o organisms (A) in a biological sample which comprises (i) dividing the sample into many parts (B); (ii) immobilizing (A) in (B); (iii) contacting (B) with at least one labeled hybridization probe (HP) and (iv) detecting bound label. Different HP (or combinations of them) are used for each (B) and HP comprise a hybridization region that is complementary to a target sequence, available for hybridization within the cell, in (A)-specific nucleic acid. The method is used to identify (A), i.e. bacteria, fungi, protozoa or multi-cellular parasites, (i) in foods (or pharmaceuticals), for process or quality control and (ii) in
                                                                                                                                                                                                                                                                                                                                                                                                                                             Detection; probe; identification; food; pathogen; tuberculos bacterium; fungus; protozoa; multi-cellular parasite; infect septic shock; necrotizing fasciitis; cystic fibrosis; mening urogenital infection; fulminant endocarditis; ophthalmitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ი
                                                                                                                                                                            Direct and rapid identification of microorganisms, useful for determining pathogens that cause fulminant infections, based hybridization with labeled immobilized probes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HONS )
                                                                                                                                                                                                                                                                                                                                                                                                                    Clostridium
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                                                                                                                                                   Claim 27; Page 29; 38pp; German.
                                                                                                                                                                                                                                                          Apfel H,
                                                                                                                                                                                                                                                                                                               17-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                  31-MAY-2001
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                                                                                                                                                                                                                                                                                                                                       17-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            intestinal bacteria using the primers shown present sequence is one of these primers.
                                                                                                                                                                                                                                                                                    (CREA-) CREATOGEN AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              perfringens
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ZH YAKULT BIOSCIENCE
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                                                                                                                                                                                                                                                          Trebesius K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention relates to the detection bacteria using the primers shown in
                                                                                                                                                                                                                                                                                                                                                                                                                   perfringens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       intestinal bacteria
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             detecting probe Cloper.
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Pred. No. 1.
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                                                                                                                                                                                                                                                            Heesemann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; DB 1;
1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and identification AAI69501-AAI69547.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tuberculosis; sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                            meningitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infection;
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RESULT
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        with infections that need very quick diagnosis, especially septic shock, necrotizing fasciltis, sepsis, exacerbation of cystic fibrosis, urogenital infections during pregnancy, fulminant endocarditis, meningitis and ophthalmitis, but also for non-fulminant infections, e.g. tuberculosis, that requires quarantining of infected subjects. The method is sensitive, specific, simple (particularly no need for lysis, culturing or microscopy), produces reliable results in typically 3 hr, is relatively inexpensive, and can identify all potential pathogens associated with a particular disease. Compared with known in situ methods, sample sizes may be 10-20 times larger, increasing sensitivity by at least an order of magnitude. The method is suitable for routine use by non-expert personnel, AAH20833-AAH20866 represent probes described in the disciplence of the invariant content of the invariant co
The invention relates to an isolated polypeptide, comprising form of mouse or human SAC1 polypeptide. The variant form is with altered preference for carbohydrates, other sweeteners of the polypeptide and its associated DNA sequence can be produced by the polypeptide and its associated polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         T 67
                                                                                                                                                                                       SAC1 polypeptide, and is associated carbohydrates or other sweeteners,
                                                                                                                                                                                          Novel isolated polypeptide comprising variant form of mouse SAC1 polypeptide, and is associated with altered preference carbohydrates or other sweeteners, useful for preventing obe
                                                                                                                                  Claim 14;
                                                                                                                                                                                                                                                                                   WPI; 2002-075162/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-APR-2000; 2000US-200794P.
28-JUL-2000; 2000US-221419P.
10-NCV-2000; 2000US-247443P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200183749-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           obesity; diabetes; transgenic embryo; blood; tongue; PCR primer; anorectic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Murine SAC1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mouse;
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                                                                                                                                                                                                                                                                                                                                                                                             WARNER
MONELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GTTGAAAGATGATGCCA 1843
                                                                                                                               Page
                                                                                                                                                                                                                                                                                                                            AA, B
Reed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GTTGAATGATGCCA 18
                                                                                                                                                                            alcoholism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene-specific oligonucleotide PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 5 A; 3 C; 6 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2001WO-US13387.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                       Beauchamp GK,
Ross D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SAC1;
                                                                                                                                                                                                                                                                                                                                                                                             LAMBERT CO.
CHEM SENSES
                                                                                                                               93; 239pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         carbohydrate; sweetener; ethanol; alcoholism; ss;
transgenic embryo; body tissue; body fluid; pancreas;
primer; anorectic; antidiabetic; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            particularly for identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.1%;
                                                                                                                                                                                                                                                                                                                                                                                               CENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.4; DB 1;
Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                            Chatterjee A,
Tordoff MG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                De
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                        be produced
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    diabetes or
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                                                                                     variant
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                                          ethanol.
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AAQ62462/c
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   screening for drugs and sweeteners. Recombinant cell lines and transgenic embryos may be used in screening for and identifying agents that induce or repress function of SAC1. Predisposition to diabetes, obesity or alcoholism can be ascertained by testing any fluid or tissue of a human (such as blood, pancreas or tongue) for sequence variations of the SAC1 gene. A sequence variation of the SAC1 locus may indicate a predisposition to diabetes, obesity and/or alcoholism and may provide a diagnostic mark. The polymucleotide can be detected in a biological sample by contacting the DNA with a probe to form a hybridisation complex sample by contacting the DNA with a probe to form a hybridisation complex
                                       primer P6(4) PTK-specific cDNA was synthesised. amplified using primers P6(4) and N5. A 2097 bp obtained. Primers B3, P12 and E6 were then used the C-terminal of the HEK2 receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              HEK-2
                                                                                                                                              Human embryonal kinase diagnosis and therapy
                                                                                                                                                                                                                                                                                                                                                                                                                    Human embryonal kinase; HEK; protein tyrosine kinase; PTK; tumour; cancer; therapy; amplification; primer; polymerase chain reaction;
 Sequence
                                                                                            RNA from human embryonic tissue was isolated.
                                                                                                                     Example 1; Page 10; 11pp; German.
                                                                                                                                                                                                                                                                       07-OCT-1992;
                                                                                                                                                                                                                                                                                                  07-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                     DE4233782-A1.
                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ62462;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ62462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                alcoholism
                                                                                                                                                                                                                 Holtrich U,
                                                                                                                                                                                                                                                                                                                           14-APR-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
                                                                                                                                                                                                                                             (CHEM-) CHEMOTHERAPEUTISCHES
                          Updated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            is then detected. The sequences represent cDNA encoding human and SAC1 polypeptides and PCR primers specific for the SCA1 genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GCTGAAGGAAGGACACA 2434
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               associated with SAC1 expression.
 B₽;
                           25-MAR-2003
                                                                                                                                                                                                                 Ruebsamen-Waigmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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(first en
                                                                                                                                                                                                                                                                       92DE-4233782
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
7 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                P6(4).
                                                                                                                                                            2-receptor
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 w
                          receptor gene.
 G; 4 T;
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                                                                                                                                                                                                                                             FORSCHUNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.4; DB 1;
Pred. No. 1.2e+02;
                                                                                                                                                                                                                 H, Strebhardt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                            protein
 0 other
                           field.)
                                                                                                                                                             1
                                                                                                                                                             useful
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                                                                                          With the use
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                                                   The cDNA was DNA fragment was in the isolation
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Matches Query Match Best Local :

Local

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Score 15.2; DB 1; Pred. No. 1.3e+02; 0; Mismatches 3;

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2323

AGTGATGTCTGGTCCTTCGG 2342

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Best Local Similarity
Matches 17; Conserv
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64159/с
                                                                                                                                                                                                                                                                                                  protein tyrosine kinase receptor. The gene encoding the receptor i related to the trk proto-oncogene. Antibodies against the encoded polypeptide are useful for diagnosis and for the treatment of tumours. The antibodies may also be radiolabelled or coupled to a cytotoxin for destruction of cancer cells. Antisense nucleic acid can be used to inhibit gene expression.

(Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tyrosine kinase; receptor; proto-oncogene; trk; detection; diagnosis; antibody; treatment; tumour; antisense; ss.
Protein kinase; treatment; disorder; cancer; human; primer;
                        Degenerate
                                                 25-MAR-2003
14-MAY-1997
                                                                                      AAT38287;
                                                                                                             AAT38287 standard; DNA; 20
                                                                                                                                                                                                                                                                            Sequence 20 BP; 6 A; 7 C; 3 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                          Three primers (AAQ64159-Q64161) were used to amplify
                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 8; 9pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      New protein tyrosine kinase and transformed cells, etc., useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1994-184380/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Holtrich U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DE4239817-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer for amplifying tyrosine kinase receptor coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
03-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ64159;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ64159 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CHEM-) CHEMOTHERAPEUTISCHES FORSCHUNG
                                                                                                                                                                                                   2323 AGTGATGTCTGGTCCTTCGG 2342
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                                                                                                                                                                            AGTGATGTCTGGACCTATGG 1
                        PCR
                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (updated)
(first entry)
                                                (updated)
(first entry)
                        primer for protein kinase cDNA isolation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92DE-4239817.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92DE-4239817.
                                                                                                                                                                                                                                        85.0%;
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                                                                                                                                                                                                                            0;
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Pred. No. 1.3e+02;
                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                      related nucleic acid - vectors, for diagnosis and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Strebhardt
                                                                                                                                                                                                                                                 Length
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                                                                                                                                                                                                                                                                                                                                                                             regions of the the receptor is
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 PCR;
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RABSULT 71
AAVO1150/c
ID AAVO11
XX AAVO11
XX AAVO11
XX AAVO11
XX PCR pr
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PN WO9731
XX WO9731
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XX 18-FEB
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             to abnormal PK activity, and cancer. Human foetal liver polypwas PCR amplified, and the products subcloned, sequenced and to produce the plasmids pLKB1-1 and pLKB1-2. When the coding region of pLKB1-1 was isolated by restriction digest, and and by northern blotting against polyA+ RNA prepared from various human organs and cultured cells, weak expression was detected almost all tissue and cell types.

(Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homeobox 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is a degenerate PCR primer for protein kinase (PK) cDNA isolation. The PK may be used to treat disorders related to abnormal PK activity, and cancer. Human feetal liver polyA+ RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA encoding protein kinase - for potential treatment of protein kinase activity related disorders and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9628554-A1
                       Brewer GJ,
                                                                                                                                                                                                                                                            28-AUG-1997
                                                                                                                                                                                                                                                                                                            WO9731012-A1
                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer; polymerase chain reaction; ampuniversal mammalian sequence tagged site;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV01150;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV01150 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 5 A; 1 C; 4 G; 4 T; 6 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example A-1; Page 23; 30pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1996-433826/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nezu J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             foetal liver; degenerate; polymerase chain reaction; ss.
                                                                             (UNIMI)
                                                                                                                                                       22-FEB-1996;
                                                                                                                                                                                                       18-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CHUS )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1813 GTGGCCGTGAAGATGTTGAA 1832
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14; Conserv
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                                                                             AIND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GTNGCNGTNAARATGYTNAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer for universal mammalian STS's.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.1%;
ilarity 70.0%;
Conservative
                             Venta PJ,
                                                                          MICHIGAN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                       96US-0012061
                                                                                                                                                                                                          97WO-US02403
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                                                                               STATE.
                             Yuzbasiyan-Gurkan V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.2; DB 1;
Pred. No. 1.3e+02;
2; Mismatches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                      reaction; amplification; UM-STS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                              genomic map;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequenced and used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                 clone; ss.
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WPI;

1997-435083/40

mammals -

useful

for

oligonucleotide primers amplifying gene regions conserved among

developing genomic maps, isolating clones and

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RESULT 72
AAVO1155
ID AAVO1
XX AVO1
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Matches
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The present sequence represents a specifically claimed oligonucleotide PCR primer. The oligonucleotide can be used for polymerase chain reaction (PCR) amplification of DNA, specifically regions of specific genes that are conserved among mammalian species, i.e. pairs of oligonucleotides from the present specification represent universal mammalian sequence-tagged site (UM-STS) primers. The primers are used to develop genomic maps, to isolate clones from libraries, to make cross-species comparisons and to develop additional genetic markers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a specifically claimed oligonucleotide PCR primer. The oligonucleotide can be used for polymerase chain reaction (PCR) amplification of DNA, specifically regions of specific genes that are conserved among mammalian species, i.e. pairs of oligonucleotides from the present specification represent universal mammalian sequence-tagged site (UM-STS) primers. The primers are used to develop genomic maps, to isolate clones from llbraries, to make cross-species comparisons and to develop additional genetic markers. UM-STS allow genomic comparisons to be made between more species.
                                                                                                                                                                                                                                                                                              New oligonuclectide primers amplifying gene regions conserved among mammals - useful for developing genomic maps, isolating clones and making cross-species comparisons
                                                                                                                                                                                                                                              Claim 1; Page 9; 26pp;
                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1997-435083/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UNMI)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV01155 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   making cross-species
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    госат
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer; polymerase chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1873 GAGATGGAGATGAAGAT 1892
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 3 A; 8 C; 2 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MICHIGAN.
MICHIGAN STATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96US-0012061
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer for universal mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.1%;
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                                                                                                                                                                                                                                              English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yuzbasiyan-Gurkan
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          site; genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amplification; UM-STS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             map; clone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STS'8.
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RESULT 73
AAZ18093
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                                     The invention provides a new method for identifying and characterising CC cells. The method for determining the genetic proximity of a first cell cand a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its can be used for determining the origin of a cell, its can individual, e.g. a fetus. They can also be used for determining the ceffect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired property. The method uses reverse transcriptase polymerase chain can be used for determining the pattern of gene expression in a selected gene family. Sequences AAZ17803-Z18342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, protein phosphatase genes, P450 mayme genes, steroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                           gene
                                                                                                                                                                                                                                                                                                                             Claim 4; Page 42; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vider B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-OCT-1998;
29-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             kinase gene;
primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genetic proximity; gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         UM-STS allow genomic comparisons to be made between
                                                                                                                                                                                                                                                                                                                                                           Identifying and gene expression
                                                                                                                                                                                                                                                                                                                                                                                                        P-PSDB; AAY14628
                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-419113/35.
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                       enes, protein phosphatase superfamily genes or cadh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCAGTCAAGTGGATGGCTCC
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97IL-0122793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98WO-IL00625
                                                                                                                                                                                                                                                                                                                                                           characterizing cells by comparing the pattern in a selected gene family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           phosphatase; P450; steroid
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Pred. No. 1.
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Best Local Similarity
                                                                                                                                                                                                                       The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is
                                                            transformed. They can be used for detecting a selected genetic defect in an individual, e.g. a fetus. They can also be used for determining the effect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired property. The method uses reverse transcriptase polymerase chain reaction (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences ARZ17803-Z18342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, expression. The gene family can be selected from a set of homeobox genes,
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                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 42; 102pp; English.
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P-PSDB; AAY14630.
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29-DEC-1997;
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                           genes, protein phosphatase genes, P450 enzyme genes, steroid or superfamily genes or cadherin superfamily genes.
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97IL-0122793.
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Pred. No. 1.
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Query Match

Sequence

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BP; 8

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other;

Best Local Similarity

1.1%; C; 5

Score 15.2; DB 1; Pred. No. 1.3e+02;

Length 20

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Query Match Best Local S Matches 17

l Similarity 17; Conserv

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DB 1;

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                                                                                                                                                                                                                                                                                                                                  The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is transformed. They can be used for detecting a selected genetic defect in an individual, e.g. a fetus. They can also be used for determining the effect of a selected treatment on a test cell. They can also be used for
                                                            obtaining cells capable of expressing an homeobox related desired property. The method uses reverse transcriptuse polymerase chain reaction (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences AAZ17803-Z18342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox gene kinase genes, protein phosphatase genes, P450 enzyme genes, steroid receptor superfamily genes or cadherin superfamily genes.
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29-DEC-1997;
                                                                                                                                                                               Sequence 20 BP; 6 A; 3 C; 4 G; 7 T; 0 other;
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P-PSDB; AAY14634.
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RESULT 78 AAZ18103

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RESULT 77
AAZ18101
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                                                                                                                                                                                 CC cell; (b) determining in the first cell and the second cell the pattern CC of expression of genes in a selected gene family; and (c) calculating a CC proximity index using a specified formula. The methods can be used for CC characterising cells, e.g. for determining the origin of a cell, its cell control of the cell; its cell caracteristic cells, whether it carries a genetic defect, or whether it is cell cransformed. They can be used for determining the cells can individual, e.g. a fetus. They can also be used for determining the cells capable of expressing an homeobox related desired containing cells capable of expressing an homeobox related desired corporary. The method uses reverse transcriptses polymerase chain cells can be used for determining the pattern of gene expression in a celected gene family. Sequences AAZ17803-Z18342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, protein phosphatase genes, P450 enzyme genes, steroid
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P-PSDB; AAY14636.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying and characterizing cells by comparing the pattern gene expression in a selected gene family
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29-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GENE-) GENENA LTD
                                                                                     Local
                                2194 AAAATAGCAGACTTTGGACT 2213
                                                                  l Similarity
17; Conserv
                                                                                                                                                                      superfamily genes
                                                                                                                                     20 BP; 6 A; 3 C; 4 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        specific
                                                                    Conservative
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97IL-0122793.
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                                                                               1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer
                                                                                                                                                                      or cadherin superfamily genes:
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                                                                                   Pred.
                                                                                     Score 15.2;
Pred. No. 1
                                                                  Mismatches
                                                                                   ..3e+02;
                                                                                                     DB 1;
                                                                                               Length
                                                                    Indels
                                                                                                       20;
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                                                                  Gaps
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RESULT 79
AAZ18105
AXU
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                                                                                                                                                                            Matches
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Best Local
                                                                                                                                                                                                                                                          receptor sur-
AAZ18105;
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                              AAZ18105 standard; DNA; 20 BP
                                                                                                                                                                                                                                         Sequence 20 BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; Page 42; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Identifying and characterizing cells by comparing the pattern gene expression in a selected gene family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene
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29-DEC-1997;
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                                                                                                                                                                            l Similarity
17; Conserv
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                                                                                                              AAAATTGCAGATTTTGGCCT 20
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97IL-0122793.
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                                                                                                                                                                                                                                         A; 3 C; 4 G; 7 T; 0 other;
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                                                                                                                                                                                          Score 15.2; DB 1;
Pred. No. 1.3e+02;
                                                                                                                                                                            Mismatches
                                                                                                                                                                     2; L.
1.3e+02;
3;
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RESULT 80
AAZ18091
ID AAZ18
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AC AAZ18
XX
AC 11-OC
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                                                                                                                                                                                                                                                                                                                                                                                                                                CC calls. The method for determining the genetic proximity of a first cell call. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a compositive for expression of genes in a selected gene family; and (c) calculating a compositive for expression of a cell, its compositive for expression and selected genetic defect in an individual, e.g. a fetus. They can also be used for determining the compositive for expressing an homeobox related desired composity. The method uses reverse transcriptase polymerase chain compositive for determining the pattern of gene expression in a compositive for determining the pattern of gene expression in a considered gene family. Sequences AAZ17803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-Z1803-
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Best Local
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29-DEC-1997;
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                  20 BP; 6 A;
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nilarity 85.0%;
Conservative
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97IL-0122793.
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                                                                                                                                                                                                                                                                                                                                                                                                    3 C; 4 G;
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                                                                                                                                                                                                                                                                                                                               Score 15.2; DB 1
Pred. No. 1.3e+02
                                                                                                                                                                                                                                                                                                                                                                                                    7 T; 0 other;
                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                         DB 1;
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11-OCT-1999 AAZ18091;

(first entry)

AAZ18091 standard;

DNA;

PTK 1

gene specific

primer.

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SXEXEXEX
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                                                                                                                                                 AAZ18185
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                                                                                                                                                                                                                                                                              Matches
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Best Local &
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genetic
                                                                                                                                                                                                                                                                                                                                                                                       can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox gukinase genes, protein phosphatase genes, P450 enzyme genes, steroid
Genetic proximity; gene expression; cell characterisation; homeobox gene
                                                                  11-OCT-1999
                                                                                                                                                                                                                                                                                                                                          Sequence
                               PTK 28 gene specific primer.
                                                                                                    AAZ18185
                                                                                                                              AAZ18185 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                      receptor superfamily genes or cadherin superfamily genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Page 42; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying and characterizing cells by comparing the pattern gene expression in a selected gene family
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29-DEC-1997;
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defect; reverse transcriptase polymerase chain reaction; RT-PCR;
gene; protein phosphatase; P450; steroid receptor; cadherin;
                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                            20 BP; 8
                                                                                                                                                                                                                                                                            Conservative
                                                                  (first
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97IL-0122793.
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                                                                 entry)
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                                                                                                                                                                                                                                                                                                                                          G; 6 T; 0
                                                                                                                                                                                                                                                                                           Score 15.2; DB 1;
Pred. No. 1.3e+02;
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                                                                                                                                                                                                                                                                                                                                          other;
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RESULT 82
AAZ18183
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC calls. The method for determining the genetic proximity of a first cell calls. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a CC proximity index using a specified formula. The methods can be used for CC characterising cells, e.g. for determining the origin of a cell, its CC genetic status, whether it carries a genetic defect, or whether it is CC transformed. They can be used for detecting a selected genetic defect in CC transformed. They can be used for detecting a selected genetic defect in cC chainfundual e.g. a fetus. They can also be used for determining the CC obtaining cells capable of expressing an homeobox related desired CC obtaining cells capable of expressing an homeobox related desired CC property. The method uses reverse transcriptase polymerase chain CC can be used in the RT-PCR reactions to determine the pattern of gene expression in a celected gene family. Sequences AAZ17803-Z18342 represent primers that CC can be used in the RT-PCR reactions to determine the pattern of gene cxpression. The gene family can be selected from a set of homeobox genes, CC kinase genes, protein phosphatase genes, P450 enzyme genes, steroid creceptor superfamily genes or cadherin superfamily genes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                Genetic proximity; gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin;
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                                                                                                                                                                                                                                                                                                                                           AAZ18183 standard; DNA;
                                                                                                                                                     PTK 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20
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29-DEC-1997;
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kinase gene; protein phosphatase; P450; steroid receptor; cadherin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying and characterizing cells by comparing the pattern
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                                                                                                                                                  specific primer.
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97IL-0122793.
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85.0%;
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Pred. No. 1
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RESULT 83
AAZ18181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of expression of genes in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is transformed. They can be used for detecting a selected genetic defect in an individual, e.g. a fetus. They can also be used for determining the cell. They can also be used for obtaining cells capable of expressing an homeobox related desired property. The method uses reverse transcriptase polymerase chain creation (RT-PCR) for determining the pattern of gene expression in a celected gene family. Sequences AAZ17803-Z18342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, protein phosphatase genes, P450 enzyme genes, steroid
                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local Similarity
                                                         Genetic proximity; gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell the pattern
Synthetic. Homo sapiens.
                                                                                                                          PTK 26 gene specific primer
                                                                                                                                                            11-OCT-1999
                                                                                                                                                                                                                        AAZ18181 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 8
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P-PSDB; AAY14718.
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29-DEC-1997;
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97IL-0122793.
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Pred. No. 1.
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XX Respi
KW Respi
KW Vacci
XX Synth
OS Chlam
XX W0992
XX W0992
XX W0992

03-JUN-1999

Chlamydia pneumoniae

Synthetic

vaccine; neutralising epitope; PCR primer; ss.

purulent

otitis media;

Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis;

erythema nodosum; pharyngitis;

PCR primer used to amplify an ORF of Chlamydia pneumoniae.

13-SEP-1999 AAX95953;

(first entry)

AAX95953 standard; DNA;

20 ВP

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                                                                                                                                                                                                         The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a cells; e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is cransformed. They can be used for determining the origin of a cell, its cell an individual, e.g. a fetus. They can also be used for determining the ceffect of a selected treatment on a test cell. They can also be used for cobtaining cells capable of expressing an homeobox related desired composerty. The method uses reverse transcriptase polymerase chain creation (RT-PCR) for determining the pattern of gene expression in a can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, prefer in homenbarase canes.
                                                                        Matches
                                                                                       Query Match
Best Local
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29-DEC-1997;
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                                                                                                                                          Sequence 20 BP; 8 A; 1 C; 5 G; 6 T; 0 other;
                                                                                                                                                                            kinase genes, protein phosphatase genes, P450 enzyme genes, steroid receptor superfamily genes or cadherin superfamily genes.
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                                    2194 AAAATAGCAGACTTTGGACT 2213
                                                                      1 Similarity
17; Conserv
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 AAAATTGGAGACTTTGGAAT 20
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97IL-0122793.
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Pred. No. 1.3e+02
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RESULT 85
AAC62208
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Matches
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                                                                      Detecting metas
or determining
  Example; Page 50;
                                                 comprises
                                                                                                                                                  WPI; 2000-687067/67
                                                                                                                                                                                                                                                                                                        13-APR-1999;
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21-NOV-1997;
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                                            metastatic potential, diagnosing metastatic prining the prognosis of a subject with prostate detecting the expression of fit-4 in a prostat
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97FR-0014673.
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78pp; English
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RESULT 86
AASO8729/c
ID AASO87
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Best Local Similarity
Matches 17; Conserva
The sequence represents a splice site within a DNA molecule encoding human PD-ATP-binding cassette (PD-ABC) protein. PD-ABC maps to chromosome 19913 3 and is expressed in various tissues including spleen, thymus, peripheral blood leukocytes, bone marrow and lymph nodes. The PD-ABC DNA molecules and proteins are used to diagnose and treat cardiovascular disorders, inflammatory disorders, dyslipidaemia, epilepsy, diseases related to abnormal calcium flux, coronary artery disease, Tangler's disease, familial high-density lipoprotein deficiency, atherosclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primers AAC62208-09 were used to amplify cDNA encoding flt-4. Flt-4 is a receptor type tyrosine kinase with 7 [g-like domains similar to other VEGF receptors. Flt-4 may play a role in lymphangioenesis. Antisense oligonucleotides can be used for detecting the metastatic potential, diagnosing metastatic prostate cancer or determining the prognosis of a subject with prostate cancer. The method comprises identifying the prostate cell in a body fluid sample and detecting the expression of flt-4 in the cell. Expression of flt-4 in a prostate cell indicates that the cell is a cancerous prostate call that has metastatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PD-ATP-binding cassette; PD-ABC; chromosome 19p13.3; spleen; thymus; ds; peripheral blood leukocyte; bone marrow; lymph node; dyslipidaemia; cardiovascular disorder; inflammatory disorder; bahormal calcium flux; epilepsy; coronary artery disease; Tangier's disease; atherosclerosis; familial high-density lipoprotein deficiency; fatty liver disease; atherosclerosis; diabetes; insulin resistance; obesity; drug screening; atherosclerosis; diabetes; insulin resistance; obesity; drug screening;
                                                                                                                                                                    Disclosure; Page 37; 77pp; English.
                                                                                                                                                                                                                        New Human PD-ABC DNA molecules and proteins for diagnosis and treatment of dyslipidaemia, epilepsy and diseases related to abnormal calcium flux
                                                                                                                                                                                                                                                                               WPI; 2001-442259/47.
                                                                                                                                                                                                                                                                                                                                                                                         24-JAN-2000;
30-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                alcoholism; retinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS08729 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PD-ABC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   potential or is a secondary tumour metastasis of a primary prostate
                                                                                                                                                                                                                                                                                                                                                       (WARN ) WARNER LAMBERT CO.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 BP; 4 A; 3 C; 9 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGAGGGATGGAGTTCCTGGC 20
                                                                                                                                                                                                                                                                                                                 Tafuri SR,
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                                                                                                                                                                                                                                                                                                                                                                                           2000US-0177889
2000US-0215405
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            form 1 DNA exon 6 5' splice site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            degeneration; hypertension; vascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       85.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.1%;
                                                                                                                                                                                                                                                                                                                 Wang
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Pred. No. 1.3e+02;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1:
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RESULT 87
AAS08820/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
      Matches
                                                                                                     19p13.3 and is expressed in various tissues including spleen, thymus, peripheral blood leukocytes, bone marrow and lymph nodes. The PD-ABC DN molecules and proteins are used to diagnose and treat cardiovascular disorders, inflammatory disorders, dyslipidaemia, epilepsy, diseases related to abnormal calcium flux, coronary artery disease, Tangier's disease, familial high-density lipoprotein deficiency, atheroscierosis, diabetes, fatty liver disease, insulin resistance, obesity, alcoholism, retinal degeneration, hypertension and vascular disease. The sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PD-ATP-binding cassette; PD-ABC; chromosome 19p13.3; spleen; thymus; ds peripheral blood leukocyte; bone marrow; lymph node; dyslipidaemia; cardiovascular disorder; inflammatory disorder; abnormal calcium flux; epilepsy; coronary artery disease; Tangier's disease; atherosclerosis; familial high-density lipoprotein deficiency; fatty liver disease; atherosclerosis; diabetes; insulin resistance; obesity; drug screening; alcoholism; retinal degeneration; hypertension; vascular disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diabetes, fatty liver disease, insulin resistance, obesity, alcoholism, retinal degeneration, hypertension and vascular disease. The sequences
                                                                 Sequence 20
                                                                                                                                                                                                                                                                            Disclosure; Page 39; 77pp; English.
                                                                                                                                                                                                                                                                                                                        New Human PD-ABC DNA molecules and proteins for of dyslipidaemia, epilepsy and diseases related
                                                                                                                                                                                                                                                                                                                                                                                                Johns MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-JAN-2000;
30-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-JAN-2001; 2001WO-US02191.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human PD-ABC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   are also used in
                                                                                                                                                                                                                                                                                                                                                                                                                             (WARN ) WARNER LAMBERT
                 Local
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                                                                                                                                                                                                                                                sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2387 CAGGGATTCCCGTGGAGGAA 2406
1.1%; cal Similarity 85.0%; 17; Conservative
                                                                                                                                                                                                                                equence represents a splice site within a DNA molecule encoding PD-ATP-binding cassette (PD-ABC) protein. PD-ABC maps to chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
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                                                                                                                                                                                                                                                                                                                                                                                                Tafuri SR,
                                                                 BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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2000US-0215405
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                                                                 A; 8 C; 3 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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                                                                                            screening assays.
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    Score 15.2; DB 1;
Pred. No. 1.3e+02;
0; Mismatches 3;
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
l.3e+02;
les 3;
                                                                                                                                                                                                                                                                                                                        diagnosis and treatment
to abnormal calcium flux
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                                Length 20;
      Indels
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                                                                      modulators of Gene 216 expression and activity. The anti-Gene 216 antibodies and antagonists may also be used to down regulate expression and activity. The anti-Gene 216 antibodies may also be used as diagnostic agents for detecting the presence of Gene 216 proteins in samples (e.g. by enzyme linked immunosorbant assay or ELISA). Disorders that may be prevented, diagnosed and/or treated by the above methods include, for example asthma, obesity and inflammatory bowel disease. The present sequence is that of a Gene 216 related primer used in examples of the invention. The primers are used in the physical mapping of the gene (ABZ72067-ABZ72088), polymorphism (SSCT) analysis (ABZ7201-ABZ72184), sequencing (ABZ72185-ABZ72268) and genotyping (ABZ72317-ABZ72362).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acids (or vectors) and proteins may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of gene 216 by expressing inactive proteins or to supplement the patients own production of Gene 216 proteins. Additionally, the nucleic acids may be used to produce the secreted Gene 216 protein, by inserting the nucleic acids into a host cell and culturing the cell to express the protein. The nucleic acids and complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acid sequences in samples and therefore which patients may be in need of restorative therapy. The Gene 216 protein may also be used as antigens in the production of antibodies against Gene 216 and in assays to identify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to isolated genes (Gene 216) from human chrome 20p13.p12 and the proteins they encode. The nucleic acids and proteins be used in the prevention, diagnosis and treatment of diseases associated with inappropriate Gene 216 expression. For example, the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; Gene 216; chromosome 20p13-p12; antiasthmatic; anorectic; antiinflammatory; gastrointestinal; gene therapy; vaccine; asthma; obesity; inflammatory bowel disease; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated genes (Gene 216) from human chromosome 20p13-p12 and proteins they encode, useful for the prevention, diagnosis and treatment of asthma, obesity and inflammatory bowel disease -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gene 216 SSCP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Page 149;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-APR-2000; 2000US-0548797
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CC Single Primer Amplification (SPA). The method comprises synthesising a CC template nucleic acid containing a predetermined sequence and hairpin CC structure with the nested oligonucleotide extension reaction. The method CC is useful for amplifying a nucleic acid, preferably for amplifying a CC family of related nucleic acid sequences to build a complex library of CC polypeptides encoded by the sequences. The engineered nucleic acid strand CC is useful for amplifying a nucleic acid strand by providing a nucleic complex library of CC acid with a predetermined sequence engineered onto its first end, a CC structure between them and contacting the engineered nucleic acid strand CC with a primer containing at least a portion of the predetermined sequence. This process is done in the presence of a polymerase and CC complementary nucleic acid strand. The method of the invention is useful CC complementary nucleic acid strand. The method of the invention is useful CC complementary nucleic acid strand. The method of the invention is useful CC complementary simultaneously more than one different target nucleic acid sequence and for amplifying simultaneously more than one different target nucleic acid complementary to the same or different nucleic acid molecules. This process is done in the presence of a polymerase and cc amplifying simultaneously more than one different target nucleic acid complementary nucleic acid sequence and for amplifying simultaneously more than one different target nucleic acid molecules. This process is accomplementary nucleic acid sequence and for amplifying an uncleic acid sequence and for amplifying simultaneously more than one different target nucleic acid molecules. This
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                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amplifying nucleic acid by synthesizing template nucleic acid containing a predetermined sequence and hairpin structure and using template for target amplification by Single Primer Amplification -
                                                                           Sequence 20 BP; 4 A; 5
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19-SEP-2001; 2001US-323400P
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                   Similarity
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   Conservative
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Score 15.2; DB 1;
Pred. No. 1.3e+02;
0; Mismatches 3
                                                                         G; 6 T; 0 other;
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Pred. No. 1.3e+02;
0; Mismatches 3
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1858 TCTGATCTGGTGTCAGAGAT 1877

Best Local Similarity

1.1%; 85.0%; w

Score 15.2; DB 1; Pred. No. 1.3e+02;

Length

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Query Match

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G; 1 T;

0 other;

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RESULT 90
ABK53119
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                     derived from gag or pol sequences, having one of the sequences fully defined in the specification, and a method for detecting HIV-1 in a biological sample, comprising mixing the sample with two or more of the amplification oligomers that specifically amplify at least one HIV-1 target sequence within gag and a pol sequence which is a protease or reverse transcriptase sequence, amplifying the target, and detecting the amplified product. The oligonucleotides of the invention may be used to diagnose HIV-1 infection. The presents sequence represents a PCR primer used to amplify the HIV-1 protease gene in the HIV detection method of the invention.
                                                                                                                                                             This invention relates to a series of nucleic acid oligomers for amplifying and detecting a nucleotide sequence of human immunodefivirus type 1 (HIV-1). The invention also comprises a labeled oligonucleotide that specifically hybridises to an HIV-1 sequence.
                                                                                                                                                                                                                                                             New nucleic acid oligomers for amplifying a nucleotide sequence from HIV-1 and probes for detecting the amplified product are specific for gag and pol regions and are useful to detect different subtypes of HIV-1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV; human immunodeficiency virus; ss; primer; gag; nrotease: reverse transcriptase; infection; PCR.
                                                                                                                                                                                                                                   Claim 1; Page 17; 37pp; English.
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(TRAN/)
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BABOLA O.
TRAN N.
VERNET G.
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/mod_base= gm
/note= "2'-O-methylguanosine"
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/mod_base=
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"OTHER= 2'-O-methyladenosine"
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RESULT 91
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              New antisense oligonucleotides targeted to nucleic acid encoding MAP/ERK kinase kinase 3 (MEKK3), useful for inhibiting the expression of MEKK3 and for treating a disease or condition associated with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; MAP/ERK kinase kinase 3; MEKK3; mitogen activated protein kinase; MAP; ERK; extracellular signal regulated kinase; infection; cytostatic; antisense therapy; tumour formation; phosphorothioate backbone;
         expression of MEKK3
                                                  WPI; 2002-329863/36.
                                                                   Ward DT,
                                                                                     (-SISI)
                                                                                                     08-SEP-2000; 2000US-0658688
                                                                                                                    07-SEP-2001; 2001WO-US28118
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RESULT 92
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to antisense oligonucleotides targetted to nucleic acids encoding micogen activated protein kinase (MAP)/extracellular signal regulated (ERK) kinase kinase 3 (MEKK3) or a splice variant of MEKK3. MEKK3 is an ubiquitously expressed serine-threonine kinase and activates only the ERK and JNK/SAPK pathways. The antisense compound is useful for inhibiting the expression of MEKK3 and for treating a disease or condition associated with the expression of MEKK3. These may also be used as research reagents and diagnostics, to distinguish between functions of various members of a biological pathway, and in the treatment of a disease or disorder, which can be treated by modulating the matter of the contract of the 
                                    New nucleic acid oligomer, useful for detecting selected regions of gag and pol genes of human immune deficiency virus, particularly for assessing drug resistance -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the expression of MEKK3. The antisense compounds are further useful prophylactically, e.g. to prevent or delay infection, inflammation or tumour formation, and as probes or primers. The present sequence is an antisense oligonucleotide targetted towards human MEKK3 cDNA.
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ss.
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                                                                                                                                                                                                                                  GEN-PROBE INC
BIOMERIEUX SA
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Pred. No. 1.3e+02;
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                       This invention relates to a novel isolated nucleic acid, gene 216, identified from human chromosome 20p13-p12. The invention also disclos regions of the 216 gene that contain single nucleotide polymorphisms (SNP's) which may be used as markers for disease susceptibility or severity. The nucleotides of the invention may have antisthmatic, antiinflammatory or anorectic activities and may be used in gene therapy. The nucleic acids, antibodies or its fragments are useful for disgnosing, preventing or treating a disorder, such as respiratory diseases (e.g. asthma, bronchial hyper-responsiveness, chronic obstructive pulmonary disease or adult respiratory distress syndrome), obesity, or inflammatory bowel syndrome. The nucleic acids are also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Keith T,
Simon J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SCHE )
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13-APR-2001;
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treating a disorder, such a
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Pandit S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ucleic acids, useful for diagnosing, preventing such as asthma, bronchial hyper-responsiveness, monary disease, obesity or inflammatory bowel
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hes 3;
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                                                                                                                                                                                            Enzymatic nucleic acid molecules having a for the treatment of arthritis, induction treatment of auto-immune diseases
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                                                                                                                                                                                                                                                                                                                                                                                                              PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           membrane; ; autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ribozyme target
                                                                                                                                                                                                                                                                                                                      McSwiggen J, Pavcc
A, Modak A, Usman
s T, Thompson JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15.2;
No. 1.
                                                                                                                                                                                                                                                                                                               Pavco P, Surgin A;
Usman N, Burgin A;
"incott F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immune response; target; cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human; rabbit; mouse; coll
arthritis; osteoarthritis;
se; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           can also be used as
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                                                                                                                                                                                                              hammer-head of graft tol
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NO:2943
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                                                                                                                                                                                                                                                                                                                                                                    Stinchcomb
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tolerance c
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residues; (ii) a 2'-C-allyl modification at position 4 of the at least ten 2'-O-methyl modifications; and (iv) a 3'-end modifications; and romelysin production the ENA's can inhibit collagenase and stromelysin production

membrane of joints for the treatment or prevention of

arthritis

(iii)

esent invention describes a novel enzymatic nucleic acid (ENA) hammerhead motif (HM) comprising: (i) at least 5 ribose es, (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii tten 2'-O-methyl modifications; and (iv) a 3'-end modification.

having

The present invention

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RESULT 95
AAX66312
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Best Local
                                                                                                                                                                                05-OCT-1995
13-DEC-1994
23-DEC-1994
23-DEC-1994
17-FEB-1995
20-APR-1995
02-MAY-1995
04-MAY-1995
07-JUL-1995
07-JUL-1995
07-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for enhancing graft tolerance or for treating autoimmune disease, and for treating allorgies and other inflammatory conditions. The ENA's can also be used in diagnosis. Albozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therappeutic treatment is lower than that required of antisense molecules, and is highly specific. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                             rheumatoid arthritis; autoimmune disease; allergy; inflammation; diagnosis; ss.
Claim 10; Page 198; 307pp; English
                                       Enzymatic nucleic acid molecules having a for the treatment of arthritis, induction
                                                                              WPI; 1996-300653/30
                                                                                                      Beigelman L, Kar
Matulic-Adamic J,
                                                                                                                                                                                                                                                                                                                                           22-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                    20-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                              WO9618736-A2
                                                                                                                                                                                                                                                                                                                                                                                                                      MUB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Arthritic condition; graft tolerance; immune response; target; cleavage; hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; ostcoarthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse B7-2 hammerhead ribozyme target SEQ ID NO:2944.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX66312;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX66312 standard; RNA; 15
                                                                                                                                                          (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1373 AGGAGATTAÇAGÇTT 1387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      h 1.1%;
Similarity 73.3%;
11; Conservative
                                                                                                                                                           RIBOZYME PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention.
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                           ę,
                                                                                                                              Gustofson J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 5
                            auto-immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                      Karpeisky A, l
c J, Jarvis T,
                                                                                                                                                                                94US-0363253.

94US-0360550.

95US-0390850.

95US-0426124.

95US-0432874.

95US-0434509.

95US-0000971.

95US-0000974.
                                                                                                                                                                                                                                                                                                                                           95WO-US15516
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                                                                                                                                                           INC
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                                                                                                   McSwiggen J, Pavco
A, Modak A, Usman
Ls T, Thompson JD,
                            diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           other;
                                                                                                                            Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                       hammer-head motif - of graft tolerance of
                                                                                                        n N, Burg
Wincott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 15;
                                                                                                        Burgin
cott F;
                                                                                                                                 Stinchcomb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                         tolerance or
                                                                                                                                 ij
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a novel enzymatic nucleic acid (ENA) CC having a hammerhead motif (HM) comprising: (i) at least 5 ribose CC residues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) CC at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification. CC The ENA's can inhibit collagenase and stromelysin production in the CC synovial membrane of joints for the treatment or prevention of arthritis, CC particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance CC in a recipient to an alloantigen of a donor. They can also be used for CC enhancing graft tolerance or for treating autoimmune disease, and for CC enhancing graft tolerance or for treating autoimmune disease, and for CC entating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of CC expression which accompany treatment with retinoids and dexamethasone. CC The concentration of ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly concentration of ribozyme required to affect a therapeutic treatment concentration of ribozyme required to affect a therapeutic treatment concentration of ribozyme required to affect a therapeutic treatment concentration of ribozyme required in the exemplification of the concentration of ribozyme required in the exemplification of the concentration.
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Best Local
                                                                                                                                        02-MAY-1995;
04-MAY-1995;
07-JUL-1995;
07-JUL-1995;
                                  Draper K,
Beigelman l
                                                                                                                                                                                                          23-DEC-1994;
17-FEB-1995;
20-APR-1995;
                   Beigelman L, I
Matulic-Adamic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rheumatoid arthritis; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Arthritic condition; graft tolerance; immune response; target; cl hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; coll stromelysin; synovial membrane; joint; arthritis; ostecarthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15
                                                                                                                                                                                                                                                            13-DEC-1994;
23-DEC-1994;
                                                                                                                                                                                                                                                                                                05-OCT-1995
                                                                                                                                                                                                                                                                                                                                  22-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                  20-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse B7-2 hammerhead ribozyme target SEQ ID NO:2945.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX66313 standard; RNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            present invention.
                                                                                       (RIBO-) RIBOZYME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    l Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 AGGAGAUUACAGCUU 15
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                                                     Gustofson J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                   Karpeisky A,
5 J, Jarvis T,
                                                                                                                                      95US-0432874.
95US-0434509.
95US-0000951.
95US-0000974.
                                                                                                                                                                                                            94US-0363253.
94US-0363254.
95US-0390850.
95US-0426124.
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94US-0354920
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                                                                                       PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 C; 4 G; 4 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     autoimmune disease; allergy; inflammation;
                McSwiggen J, Pavcc
A, Modak A, Usman
is T, Thompson JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4
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Pred. No. 97;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immune response; target; cleavage;
                   Pavco P, Stinchco Usman N, Burgin A; JD, Wincott F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
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                                                       Stinchcomb DT;
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WPI; 1996-300653/30

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RESULT 97
AAP52668
ID AAP5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; cardinit; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBD3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
                                   WPI; 2001-041421/05
                                                                                                                                                                                                                                            21-JUN-1999;
                                                                                                                                                                                                                                                                                                       21-JUN-2000; 2000WO-AU00693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF52668;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF52668 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200078341-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IGF-I oligonucleotide #3628.
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                                                                                                                                                                         MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    treatment of arthritis,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 4 A; 3 C; 4 G; 4 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                      Werther
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                      Edmondson
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Pred. No. 97;
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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an CC antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 CC receptor, IGF binding protein [IGFBP]-2 or IGFBP3], which is capable of CC inhibiting or reducing growth factor mediated cell proliferation, CC oligonucleotide which can be used to design the antisense CC oligonucleotide which can be used to design the antisense CC oligonucleotides of the present invention (see AAF45151 and CC oligonucleotides of the present invention (see AAF45151 and CC psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, CC keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the CC skin, a hypermeovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other CC sclerotic disease, kidney disease, hyperproliferation of the inside of the other throat the present of the inside of the other westels or any other hypermediated malignancies, other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisonucleic acid that inhibits or reduces growth factor mediated cell
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                proliferation and/or inflammation -
                                                   blood vessels or any other hyperplasia.
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BP; 0
Α,
4 C;
6 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      an antisense
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밁 ş Matches Query Match Best Local (2329 l Similarity 15; Conserv GTCTGGTCCTTCGGG GTCTGGTCCTTCGGG Conservative 100.0%; 15 0; Score 15; Pred. No. Mismatches DB 1; 0 Length 15; Indels ٥, Gaps

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RESULT 98 AAF52669 Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor, 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrheea; ruba; keratosis; neoplasia; seleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; AAF52669 standard; DNA; IGF-I oligonucleotide #3629. neovascular condition of the (first entry)

28-DEC-2000.

21-JUN-2000; 2000WO-AU00693

21-JUN-1999; 99US-0140345.

(MURD-) MURDOCH CHILDRENS RES INST.

Wraight CJ, Werther GA, Edmondson SR

WPI; 2001-041421/05

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an a nucleic acid that inhibits or reduces growth factor mediated an antisense

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RESULT 99
AAF52670
ID AAF52670
ID AAF52670
AAF52
XX AAF52
XX AAF52
XX Antis
KW Cytos
KW Cytos
KW Skin
KW Hypex
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatclogical; cardiant; virucide; opthhalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
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                                                                                                                                       Ameliorating the effects of a disorder, e.g. psoriasis, ladministering UV (ultra-violet) treatment (optional) and nucleic acid that inhibits or reduces growth factor media
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97;
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The present invention relates to a method for ameliorating the effects

Example

8; Page 84; 201pp; English

exocrine cells. The invention is useful in generating insulin-secreting cells from precursor stem cells or adult pancreatic exocrine cells is useful for generating glucose sensitive insulin secreting beta cells suitable for transplantation, and for in situ development of insulinsecreting cells in a patient. The method is also useful for preventing

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RESULT 100
AAD47315/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP]), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hypernevoascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other scleroid consense, kidney disease, hyperproliferation of the inside of blood morels of the inside of
                                                                       The invention relates to a method for generating insulin-secreting cells from precursor stem cells or adult pancreatic exocrine cells. The method comprises exposing the precursor exocrine cells to: a nucleic acid molecule encoding neurogenin 3 (ngn3) or NeuroD/beta2; or an activator of ngn3 or NeuroD/beta2 gene expression, under conditions effective to generate the insulin-generating cells from the precursor or
                                                                                                                                                                                                                                      Generating insulin-secreting cells from precursor stem cells or adult pancreatic exocrine cells, for generating glucose sensitive insulin secreting beta cells for transplantation, comprises using neurogenin3
                                                                                                                                                                                                                       secreting beta cells or NeuroD/beta2 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; insulin-secreting cell; neurogenin 3; ngn3;
pancreatic exocrine cell; transplantation; RT-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAD47315 standard; DNA;
                                                                                                                                                                                           Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human RT-PCR reverse primer for synaptophysin DNA isolation
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                                                                                                                                                                                                                                                                                                                                      Heimberg H,
                                                                                                                                                                                           Page 36;
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97;
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밁 S

(as in cancer therapy)

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Primers AAV60743-V60745 were used to PCR amplify codons 1-163 of the chuman cyclin-dependent kinase 4 (hCDR4). The amplified product was used to generate a fusion protein comprising part of the hCDR4 sequence linked to codons 154-302 of the yeast PHO85 gene. The fusion protein is used to screen for compounds that affect mammalian cell cycle regulatory proteins. The method comprises administering a compound to a cell line, which contains a reporter gene linked to an Upstream Activation Sequence (UAS) and a promoter, where the UAS binds a transcription control factor (TCF) which is regulated through cyclin-dependent kinase (CDK) phosphorylation. Also included in the construct is an effector gene providing a gene product to permit normal cyclin/CDK regulation of the TCF. Expression of the reporter gene is then analysed in the cell line, thereby determining whether the compound affects the normal regulation. The method can be used to identify inhibitors and activators of mammalian cell cycle regulatory proteins, especially inhibitors and activators of cyclins, CDKs, cyclin/CDK complexes, cyclin kinase inhibitors (CKIs), and cyclin/CDK/CKI complexes. The identified agents can be used for stimulating growth of cells (as in wound healing), or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                    Screening for agents that effect cell cycle regulatory proteins using a cell line that expresses a reporter gene in response to regulation through phosphorylation by a cyclin/CDK system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer; amplification; yeast; UAS; upstream activating sequence; transcription terminator; cell cycle; Upstream Activation Sequence; UAS; promoter; phosphorylation; cyclin; cyclin-dependent kinase; CDK; vector; cyclin kinase inhibitor; CKI; growth; wound healing; cancer therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  premature differentiation of precursor stem cells into insulin-secreting beta cells and for identifying compounds that prevent or activate beta cell differentiation. The present sequence is human RT-PCR primer for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                    Example 4; Page 75;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-251302/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bitter GA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer #2 for human CDK4 codons 1-163.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (BITT-) BITTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-OCT-1996;
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96US-0029127.
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                                                                                                                                                                                                                                                                                                                                                                  93pp;
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100.0%; Pred. No.
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                                                                                                                                                                      modulate expression of G-alpha-16. G-alpha-16 is a human G protein which interacts differentially with several receptor types including members of the opioid and chemokine receptor families. A series of antisense oligonucleotides have been designed to target different regions of the human G-alpha-16 RNA. They may be used to inhibit the expression of G-alpha-16 in human cells and tissues and thus to treat diseases associated with G-alpha-16, sue he prepared thus to treat diseases, especially cancer. Infections, inflammation or tumour formation can be prevented or delayed. The compounds can be used in research and disgnostics in sandwich and other assays.
                                                                                                                                                                                                                                                                                                                         The
                                                                                                                 Note: The sequence has a phosphorothioate backbone and may be either an oligodeoxymucleotide or a chimeric oligonucleotide containing 2'-methoxyethyl (2'-MB) wings and a deoxy gap. Toumber given above corresponds to the oligodeoxynucleotide s
                                                                                                                                                                                                                                                                                                                                                                                A new antisense compound for inhibiting the expression of human G-alpha-16 and treating, preventing or delaying infections, inflammation or hyperproliferative disorders such as cancer -
                                                                                       Sequence 18
                                                                                                                                                                                                                                                                                                                                                      Example 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-412354/35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; G-alpha-16; G protein; cytostatic; hyperproliferative disorder; cancer; inflammation; infection; antisense inhibition; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human G-alpha-16 antisense oligonucleotide ISIS# 20849
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    regulating excessive cell growth and division
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS
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                                                                                                                                                                                                                                                                                                                         present sequence is an antisense oligonucleotide used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2203 GACTTTGGACTCGCCAGA
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                              16;
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16; Conserv
                                             Similarity
 CCAGCGCCTGGAAGAGAA 1371
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                                                                                       B₽;
                                                                                                                                                                                                                                                                                                                                                     Page 73; 100pp; English.
                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                      7 C; 4 G; 5 T;
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Pred. No. 1.
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                                                                                                                                                                         with inducible nitric oxide synthase, e.g. diabetes, immunological disorder, cardiovascular disorder, neurological disorder or ischaemia/reperfusion injury. The antisense oligonucleotides are also useful for research and diagnostics. The present sequence is that of an antisense 2'-O-methoxyethyl gapmer oligonucleotide with a phosphorothicate backbone, a central "gap" region of ten nucleotides flanked by four nucleotide 2'-MOE (2'-methoxyethyl) wings and 5-methylcytidine residues throughout the oligonucleotide. The antisense oligonucleotide is targeted to human inducible nitric oxide synthase (NOS)
                                                                                                                                                                                                                                                                                                                                                                                                                                          expression of inducible nitric oxide synthase. The antisense compounds have immunomodulator, antidiabetic, cardiovascular, cardiant, neuroprotective, disorder and vasotropic activity. The antisense oligonucleotides are useful for inhibiting the expression of inducible nitric oxide synthase in cells or tissues. In particular, the antisense oligonucleotides are useful for treating diseases or disorders associated with inducible nitric oxide synthase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modulate expression; immunomodulator; antidiabetic; cardiovascular; cardiant; neuroprotective; vasotropic; ischaemia; reperfusion injury; 2'-O-methoxyethyl; phosphorothioate; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to antisense compounds, especially oligonucleotides, which are targeted to a nucleic acid encoding inducible nitric oxide synthase and which specifically hybridise to and modulate expression of inducible nitric oxide synthase. The antisense compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense oligonucleotides for modulating the expression of inducible nitric oxide synthase in cells or tissues, particularly useful for treating e.g. immunological, cardiovascular or neurological disorders, or ischaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human inducible NOS antisense oligonucleotide SEQ
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                                                                                         Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 85; 144pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dean NM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cowsert LM;
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                                                                                         G; 7 T; 0
Score 14.8; DB 1;
Pred. No. 1.3e+02;
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Best Local Similarity

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                                                                             targeted to a nucleic acid encoding Her-4 (a receptor/tyrosine kinase) The antisense oligonucleotides are used to inhibit the expression of Her-4 in cells or tissues in vitro. They can be used in diagnostics, therapeutics, prophylaxis and as a probe in research reagents. The antisense oligonucleotides can be used to prevent or delay infection, inflammation or tumour formation. AAH25315-AAH25398 represent antisense oligonucleotides which are received.
                                                                                                                                                                                                                                                                                                                                                 Compound for inhibiting the expression of Her-4 (a receptor/tyrosine kinase) e.g. in preventing tumour formation, comprises an antisense oligonucleotide that hybridizes to a nucleic acid encoding Her-4 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified_base
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                                                                                                                                                                                                                                                                                                  Example 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-388929/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-JUL-2000; 2000US-0632580.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-JUL-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense oligonucleotide targeted to human Her-4 coding region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammation; tumour; phosphorothioate;
  BP;
                                                                                                                                                                                                                                                                                                Column 45-46; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cowsert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000US-0632580.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PHARM
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  4 A;
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15..18
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  7 C; 0
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"2'-methoxyethyl (2'-MOE) nucleotides"
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"2'-deoxynucleotides'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cytidine residues are 5-methylcytidines"
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  G; 7 T; 0 other;
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RESULT 106
AAA85912
ID AAA859
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AC AAA859
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DT 04-DEC
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Best Local S
                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                              Representative examples of ribozyme recognition sites are given in AAAB2415 to AAAB6787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells.
                                                                                                                                                                                                                                                                                              The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent konter than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1.
                                04-DEC-2000
                                                                                                                                                                                                                    Sequence 19
                                                                                                                                                                                                                                                                                                                                                                New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1
                                                                                                                                                                                                                                                                                                                                           Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Tritz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ribozyme; hairpin; hammerhead; gene therapy; vasotropic;
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                                                                         AAA85912 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-412314/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (IMMU-) IMMUSOL INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         restenosis;
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           25
                                                                                                                                                  1320 GATATCCTTTCACTCTGC
                                                                                                                                                                                                                                                     ribozyme is resistant to endonuclease activity and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1884 GATGAAGATGATTGGGAA 1901
                                                                                                                                                                           al Similarity
16; Conserv
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          hs ribozyme binding site #20.
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                                                                                                                                 GATTTCCTTTCATTCTGC
                                                                                                                                                                                                                                            in
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Welch PJ,
                                                                                                                                                                                                                     BP; 3 A;
                                                                                                                                                                                                                                                                                                                                           Page 99; 109pp; English.
                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                           restenosis treatment.
                              (first entry)
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                                                                                                                                                                                    1.1%;
                                                                                                                                                                                                                     5 C; 2
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                                                                           ВP
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                                                                                                                                                                                                                     G; 9 T; 0 other;
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                                                                                                                                                                                    Score 14.8; DB
Pred. No. 1.4e+(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. 1.3e+02;
0; Mismatches 2;
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                                                                                                                                                                                    DB 1;
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                                                                                                                                                                                              Length 19;
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03-MAY-2001 WO200130362-A2 Synthetic Homo

sapiens

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RESULT 107
AAH61073
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Best Local :
                                                                                                                       Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerar; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostat; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDKI, PCNA and Cyclin B1. Representative examples of ribozyme cognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is
                                                                                                                 sickle
                                                                                                                                                                                                                                                                       Cdc25 hs ribozyme binding site SEQ ID NO:3497.
                                                                                                                                                                                                                                                                                                                                                            AAH61073 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 99; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New hairpin and hammerhead ribozyme RNA encoding a cyclin or cell-cycle PCNA and Cyclin B1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                                                                                                                                                                                                                                                                     10-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tritz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       1320 GATATCCTTTCACTCTGC 1337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
                                                                                                                 cell
                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GATTTCCTTTCATTCTGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             yme is resistant to endonuclease activity in restenosis treatment.
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                                                                                                                retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                             ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.8;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.4e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         other than CDK1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                             vulnerary;
                                                                                                                                                                                      cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cleaves
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local (
                                                                                                                                                                                                                                                           Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation, cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipporiatic; dermatological; antiseborrheic; antidiabetic; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH57577 to AAH62099 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can be used for the state of the contract of the cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammat. matrix metalloproteinases, growth factors and cell-cycle dependent
                                                                                                                                                                                                        antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cdc25 hs ribozyme binding site SEQ ID NO:3498.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (I) which cleaves RNA encoding a cytokine involved in
      WO200130362-A2
                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-300427/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-OCT-2000; 2000WO-US29500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAH61074 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (IMMU-) IMMUSOL INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1320 GATATCCTTTCACTCTGC 1337
                                                                                                                                                                                  cel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N
                                                                                                                                                    cell retinopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                               carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A; 5 C; 2 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tritz R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0161532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.8; DB 1;
Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                            wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ฌ
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RESULT 109
AAQ49761
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cc ribbzyme (I) which cleaves RNA encoding a cytokine involved in clinical control of the control of the control of the cyclin, cell-cycle control of the cyclin, cyclin, cell-cycle control of the cycle cycl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammat matrix metalloproteinases, growth factors and cell-cycle dependent
                                                                                                                                                                                                                            MIS; Mullerian Inhibitory Substance; transforming growth factor; inhibin;
                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003
20-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ49761 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 2 A; 5 C; 2 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           scar. AAH57577 to AAH62099 represent sequences exemplification of the present invention.
                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                            Membrane serine/threonine kinase receptor DNA primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 326; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-300427/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Robbins JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-OCT-2000; 2000WO-US29500.
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                                                                                                                                                                                                          oone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity hes 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1320 GATATCCTTTCACTCTGC 1337
                                                                                                                                                                          polymerase chain reaction; primer; amplification; ss.
                                                                                                                                                                                                    sforming growth factor; inhibin; BMP;
morphogenesis protein; serine/threonine kinase receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or eye disease and scarring.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GATTICCTTTCATTCTGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention describes a method for treating a proliferative disease and scarring. The method involves administering a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tritz R
                                                                                                                                                                                                                                                                                                                                                         (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0161532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.8; DB 1;
Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                          receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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30-SEP-1993

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RESULT 110
AAT36072
ID AAT360
XX AAT360
XX AAT360
XX 25-MAR
DT 25-MAR
DT 30-OCT
XX Transf
XX Muller
XX Wound
KW Wound
KW Wound
KW Wound
XX Synthe
PN US5538
XX US5538

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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                         04-NOV-1993;
18-MAR-1992;
11-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The primers given in AAQ49761 and AAQ49762 were used in the isolation of four novel membrane serine/threonline kinase receptor cDNAs.

Misrl (AAQ49763) is believed to encode an isoform of the rat
MIS receptor. Misr2A/misr2B (AAQ49764), misr3 (AAQ49765) and misr4
(AAQ49766) are believed to encode monomeric isoforms of the rat
inhibin receptor and/or BMP receptor.

(Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transforming growth factor beta type I recept wound healing; tumour treatment; rat inhibin;
New isolated TGF-beta type I receptor DNA - used to develop prods for diagnosis and therapy, e.g. for treating tumours or promoting
                                                                                                                                                                                     (GEHO )
                                                                                                                                                                                                                                                                                                                                                              04-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mullerian inhibiting substance receptor; MISR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Transforming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 23; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New receptors of the transforming growth factor-beta receptor family - comprising Mullerian Inhibitory Substance Receptors and
                                                                                   WPI; 1996-353830/35
                                                                                                                                                                                                                                                                                                                                                                                                                23-JUL-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polymerase chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT36072 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inhibin receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1993-320743/40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-MAR-1992;
11-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1813 GTGGCCGTGAAGATGTT 1829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -
                                                                                                                                                                                     GEN HOSPITAL CORP
UNIV DUKE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.0%;
Similarity 82.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GTGGCCGTSAARATYTT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                    Gustafson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      growth factor beta receptor superfamily PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (updated)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gustafson
                                                                                                                                                                                                                                                         93US-0149105.
92US-0853396.
93US-0029673.
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93US-0029673.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; 2 C; 4 G; 5 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                    Σ
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                                                                                                                                    Hе
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.6; DB 1;
Pred. No. 1.3e+02;
                                                                                                                                    Ξ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ¥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             receptor; gene
                                                                                                                                    ×
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGF-beta receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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RESULT 111
AAH40382/c
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Degenerate PCR primers were designed based on two highly conserved regions within the cDNA encoding a murine activin receptor, human and porcine TGF-beta type II receptor and the daf-1 receptor of C.elegans. The primers (see AAT36072 and AAT36073) were used for amplifying clones present in a 14.5 day foetal rat urogenital ridge cDNA COS cell expression library. Four clones encoding portions of four novel polypeptides (all putative serine/threonine kinases) were obtained and designated pGEM7-Misr1, 2, 3 and 4. The inserts from these clones were used as probes to isolate full-length cDNA sequences for each of the four TGF-beta type I receptors, Misr1 is believed to encode an isoform of the rat Mullerian Inhibiting substance (MIS) receptor, while misr2A/misr2B, misr3 and misr4 are
                                                                                                                                                                                                                                                                                                                                                                                                                                                Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                               Claim 1;
                                                                                                                New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                               Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                     26-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP specific lower PCR primer SEQ ID 3178.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-AUG-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH40382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 3 A; 2 C; 4 G; 5 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            believed to encode monomeric isoforms of the rat inhibin receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Column 14; 44pp; English
                                                                                                                                                                                                                                                                   15-OCT-1999;
                                                                                                                                                                                                                                                                                                  13-OCT-2000; 2000WO-US28436
                                                                                                                                                                                                                                                                                                                                                                    WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                      inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and/or BMP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             wõund healing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1813 GTGGCCGTGAAGATGTT 1829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GTGGCCGTSAARATYTT 17
                                                                                                                                                                                                                                ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                               Page 66; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                      forensic
                                                                                                                                                                                                                                                                   99US-0160096.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                    investigation; paternity analysis; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.6; DB 1;
Pred. No. 1.3e+02;
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                                                                                                                a nucleic
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention

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ARTSSILT 112
AATTS623/C
ID AATTS62
XX AATTS62
AC AATTS62
XX 25-MAR
DT 25-MAR
DT 16-APR
XX PRODUCT
KW disord
KW PRODUCT
KW disord
KW WPSYSTEM
KW MULTIP
KW WO9624
XX SYNTHE
PN WO9624
XX 15-AUG
PP 05-PEB
PR 09-PEB
PR 18-SEP
PR WPI; 1
XX REPTIONS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC includes kits for determining the presence or absence of a SNP, using the CC oligonuclectides of the invention. The PCR primers are used to amplify a CSNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonuclectides are useful for genotyping a nucleic acid sample by performing a single-nuclectide primer extension reaction. The CC oligonuclectides are useful for determining the presence, absence or cc identity of a SNP and for genotyping nucleic acid samples, for e.g. to consider the presence of the presence of the individual or group of conditional properties and for genotyping nucleic acid samples, for e.g. to consess by association analysis the genotype of an individual or group of conditional properties. The presence of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                   Oligo:nucleotide which reduces CD28 gene expression for treating immune system diseases, e.g. graft vs. septic shock, psoriasis, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reduction; T cell; CD28; gene expression; treatment; immune system; disorder; graft versus host disease; septic shock; viral disease; psoriasis; type I diabetes mellitus; thyroiditis; sarcoides;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     multiple sclerosis; uveitis; rheumatoid arthritis; systemic lupus erythematosus; inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                                09-FEB-1995;
18-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                        05-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                        15-AUG-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT36213 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18
                                                                                                                       WPI; 1996-384228/38.
                                                                                                                                                                       Tam RC
                                                                                                                                                                                                                      (ICNC ) ICN
                                                                                                                                                                                                                                                                      18-SEP-1995;
                                                                                                                                                                                                                                                                                            09-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9624380-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a human SNP containing DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1382 CAGCTTCCCCAGACT 1396
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligo targetting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligonucleotide;
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                                                                                                                                                                                                                      PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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(first entry)
                                                                                                                                                                                                                                                                                          95US-0387041.
95US-0529878.
95US-0387041.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CD28 stop codon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ΡP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.6; DB 1;
Pred. No. 1.4e+02;
l; Mismatches 0;
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Ti
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AAX9037/C
ID AAX903
XX AAX903
XX AAX903
XX AAX903
XX CD28,
XW CD28,
XW IMMUNE
XX Synthe
XX Synthe
XX IS-SEP
XX IS-SEP
XX IS-SEP
XX IS-SEP
XX ITAME/
XX TAME/
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Query Match
Best Local S
Matches 15
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Best Local
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                                                                                                                                                       The present invention describes a method for inhibiting the expression of CD28, IL-2, gamma-interferon or IL-8 in a mammal. The method comprises subcutaneous administration of an oligonucleotide (OGN). AAX90288 to AAX90291 represent specifically claimed OGNs for use in the method. The OGNs are used for the treatment of immune system-mediated diseases. AAX90292 to AAX90323 represent oligonucleotides used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX90307 standard; DNA; 16
                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                  Treatment of immune system-mediated diseases by inhibiting expression of CD28, IL-2, gamma-interferon or IL-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CD28; inhibition; antisense oligonucleotide; interleukin 2; immune system mediated disease; gamma-interferon; IL-8; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide RT14 used in an Example from US5932556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX90307;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 16
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                                                                                                                                                                                                                                                                                                                                                                                  Example; Column 15; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-443609/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-SEP-1999
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  l Similarity
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В
С
                                                                                                          BP;
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  Conservative
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                                                                                                          1 A;
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                                                                                                          5 C;
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                         1.0%;
                                                                                                             6 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6
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Score 14.4; DB 1
Pred. No. 1.3e+02
0; Mismatches
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Pred. No. 1
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                                                                                                             0 other;
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                                                     DB 1;
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                                                     Length 16;
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                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                        cervical and ovarian cancers and to make hormonal drugs for retarding the conversion of a hormone-dependent cancer into a hormone-independent cancer. The drug can retard the change of hormone-dependent cancers into hormone-independent cancers effectively. The present sequence represents a PCR primer which is used in an example from the present
Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hormonal drug-containing agents for retarding conversion of hormone-dependent cancers into hormone-independent cancers, useful e.g. for treating prostate and breast cancers
                                      IL-1 beta hairpin/hammerhead ribozyme recognition
                                                                                           AAH61977
                                                                                                                                                                                                                                                                                       Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a hormonal drug-containing agent (I) for retarding the conversion of a hormone-dependent cancer into a hormone-independent cancer. The agents can be used to treat prostate, breast,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hormone dependent cancer; hormone independent cancer; hormonal drug; prostate cancer; breast cancer; cervical cancer; ovarian cancer;
                                                                 10-SEP-2001
                                                                                                                    AAH61977 standard; DNA; 16 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 19; 31pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matsutani E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-APR-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human fibroblast growth factor antisense PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA29601;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (TAKE ) TAKEDA CHEM IND LTD.
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                                                                                                                                                                                                           2323 AGTGATGTCTGGTCCT 2338
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                                                                                                                                                                                                                                                                                       BP; 4
                                                                                                                                                                                                                                     Conservative
                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98JP-0286793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-JP05533.
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                                                                                                                                                                                                                                                                                     C; 1 G; 2 T; 4 other;
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                                                                                                                                                                                                                                                Score 14.4; DB 1;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                          Length 16;
                                        Bite
                                                                                                                                                                                                                                    Indels
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                                        NO:4401.
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RESULT 116
AAX72965
ID AAX729
XX
AC AAX729
XX
AC AAX729
XX
DT 28-JUL
XX
DE Mouse

AAX72965

standard;

RNA;

17

28-JUL-1999

(first entry)

Mouse flk-1 VEGF receptor hammerhead ribozyme substrate #398.

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Matches Query Match Best Local

15; Conser

Conservative

0;

1.0%;

Score 14.4; DB 1; Pred. No. 1.3e+02; Mismatches

DB 1;

Length 16; Indels

0,

Gaps

0

2324 GTGATGTCTGGTCCTT 2339

GTGATGTCTGGTCCAT 16

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cleaves RNA encoding cytokine involved in inflammation. (1) can be used in gene therapy. (1) and (11) are useful for treating proliferative skin diseases such as pooriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scarr. AAH57577 to AAH62099 represent sequences used in the
                                                                                                                                                                                                                                                                                                     ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering nucleic acid molecule (II) comprising a promoter operably linked nucleic acid segment encoding (I). (I) can have antipsoriatic, nucleic acid segment encoding (I). (I) can have antipsoriatic,
                                                                                                                                                                                                                                                                                                                                                                                                                      skin
Sequence 16 BP; 2 A; 3 C; 5 G; 6 T; 0 other;
                                                                                                                                                                                                                                                              dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treating proliferative skin or eye diseases and scarring, ribozymes that cleave RNA encoding cytokines involved in matrix metalloproteinases, growth factors and cell-cycle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antisickling; ophthalmological; keratolytic; gene therapy; viral wart;
atopic dermatitis; actinic keratosis; squamous cell carcinoma;
basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostat antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide;
                                             exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 20; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-300427/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Robbins JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                      or eye disease and scarring.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tritz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0161532.
                                           the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                    describes a method for treating a proliferative nd scarring. The method involves administering a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic;
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ID AAALISS
XX AAALISS
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XX AAALISS
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (MgF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour
                              myopic degeneration; psoriasis; verruca vilgaris; angiofibroma;
tuberous sclerosis; pot-wine stain; Sturge Weber syndrome;
Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss
                                                                                                                        ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma;
                                                                                                                                                                                          Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; anglogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vascular endothelial growth factor receptor; VBGF receptor; flt-1; VlsC, kDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid archritis; ocular disease; fms-like, tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAY-1997.
                                                                                                                                                                                                                                                                                                                                                           Human TIE-2 substrate sequence SEQ ID NO:2114.
                                                                                                                                                                                                                                                                                                                                                                                                                              19-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA18888 standard; RNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present invention describes nucleic acid molecules which modulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1822 AAGATGTTGAAAGATG 1837
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12; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ŗ,
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                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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95US-0005974.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         В₽
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.4; DB
Pred. No. 1.4e+(
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Stinchcomb D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4e+02;
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RESULT 118
ABN08382/c
ID ABN083
XX
AC ABN083
XX
TY
AC ABN083
XX
DT 29-MAY
XX
Human
XX
KW Human;
KW Human;
KW muscle
KW skelet
XX
OS Homo s

29-MAY-2002

(first entry)

ABN08382 standard; DNA; 17

Human GDMLP-1 17-mer scanning

SEQ ID

NO:5

sequence SEQ

hGDMLP-1; heart; heart disease; ID NO:8374.

Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens

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Matches

Similarity 9; Conserv

Conservative

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Gaps

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AGTGATGTCTGGTCCT 2338

AGUGAUGUAUGGUCCU 16

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The present invention describes enzymatic nucleic acid molecules with CC RNA cleaving activity, which specifically cleave RNA encoded by an aryl CC hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 CC gene, an integrin alpha 6 subunit gene, or a fie-2 gene. AAA16775 to AAA17561 to AAA17622 represent ribozyme sequences for ARNT, CC aAA17168 to AAA17560 and AAA17625 to AAA17684 represent their CC corresponding target sequences, AAA17685 to AAA18385 and AAA19087 to AAA19154 represent ribozyme sequences for Tie-2, and AAA19386 to AAA19086 CC AAA19155 to AAA19222 represent their corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to CC aAA19232 to AAA21961 and AAA21501 to AAA21555 represent ribozyme cc sequences for integrin alpha 6 subunit, and AAA20362 to AAA21500 and AAA21505 to AAA21509 to AAA21688 represent their corresponding target sequences; CC AAA16889 to AAA22169 and AAA21501 to AAA21552 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA23262, AAA23343 to AAA23422 represent their corresponding target sequences. The ribozyme of the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, creat cancer, diabetic retinopathy, age related ce specially used to treat cancer, diabetic retinopathy, age related ce specially used to treat cancer, diabetic retinopathy, age related consovascular glaucoma, myopic degeneration, sontasis, verruca vulgaris, and other syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2, integrin subunit bara-3 or integrin subunit bara-3 or integrin subunit bara-3 or integrin subunit bara-3 or integrin subunit bara-6 or integrin subunit bara-8 or integrin subunit 
                      Query Match
Best Local :
                                                                                          Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors - \,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 56; Page 123; 305pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC
                                                                                                                                            subunit
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                                                                                          BP; 4 A;
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                                                                                                                                          alpha-6,
                    1.0%;
                                                                                          2 C;
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                                                                                             5 G;
                                                                                                                                          or integrin subunit
Score 14.4; D
Pred. No. 1.4e
6; Mismatches
                                                                                             6 U; 0 other;
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                                             DB 1;
                                                                                                                                               beta-3.
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                                             Length 17;
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                                                          Matches
                                                                                       Query Match
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30-JAN-2001
30-JAN-2001
                                                                                                                                                                                                             concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-MAY-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                                                                                     N.B. The sequence data for this patent did not form precification, but was obtained in electronic format at ftp.wipo.int/pub/published_pct_sequence.
                                                                                                                                                                                                                                                                                                                                                          hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a human genome-derived myosin-lil protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The
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                                                                                                                        Sequence 17
                                                                                                                                                                                                   invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    myosin-like
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                                                                            Local
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                             1506 CCAGCCGGCTGTGCAC 1521
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                                                            Conservative
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2001WO-US00669.
2001WO-US00670.
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2001WO-US00667.
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                                                                                                                      4 A;
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                                                                         1.0%;
                                                                                                                      5 C; 6 G; 2 T; 0 other;
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                                                                         Score 14.4; DB 1;
Pred. No. 1.4e+02;
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                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human genome-derived myosin-like
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                                                                                      Length 17;
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t directly from WIPO
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                                                          Gaps
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Human GDMLP-1 17-mer scanning
                                                                                           ABN08383 standard; DNA;
                            (first entry)
                                                                                            17
 SEQ
ID NO:5 sequence SEQ
 ID NO:8375
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muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss. genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; heart disease;

Homo sapiens.

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26-MAY-2000;
21-SEP-2000;
                                                                                                       30-JAN-2001;
                                                                                                                                            30-JAN-2001;
                                                                                                                                                            30-JAN-2001;
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                                                                                                                                                                                                                                                                                               WO200192524-A2
 (AEOM-)
                                       30-JAN-2001;
30-JAN-2001;
                                                                 30-JAN-2001;
                                                                              30-JAN-2001;
                                                                                          30-JAN-2001;
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                                                                                                                                 30-JAN-2001;
AEOMICA INC.
                       2001WO-US00669.
2001WO-US00670.
2001US-266860P.
                                                              2001WO-US00662.
2001WO-US00663.
2001WO-US00664.
2001WO-US00665.
2001WO-US00666.
2001WO-US00667.
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۲ Ji Y, Penn SG, Hanzel DK, Rank DR, Chen ξ Shannon ME

2002-179446/23.

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1 -

Disclosure; SEQ ID 8375; 214pp; English

CC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 can be used is gene therapy and vaccine production. The CC and quantify hGDMLP-1 nucleic acids in samples, as amplification consists that it is a substrates for the recombinant engineering CC and quantify hGDMLP-1 nucleic acids in samples, as amplification consists the protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may CC be used as immunogens to raise antibodies that specifically recognise CC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific CC deficiency in hGDMLP-1 production, and in vaccines or for replacement CC diagnosing a disorder associated with the expression of hGDMLP-1 in CC diagnosing a disorder associated with the expression of hGDMLP-1, in CC chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present convertion. The present invention. it invention describes a human genome-derived myosin-like (hGDMLP-1). The protein and polynucleotide sequences of

N.B. The sequence data for this patent did not specification, but was obtained in electronic at ftp.wipo.int/pub/published_pct_sequence format form part of the printed

RESULT 119 ABN08383/c

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RESULT 120
ABN09009
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30-JAN-2001;
30-JAN-2001;
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21-SEP-2000;
27-SEP-2000;
The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynuclectide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific
                                                                                                                                                                               New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser description ionization, comprises human myosin-like protein hGDMLP-1 -
                                                                                                                                                                                                                                              WPI;
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30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            muscle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; genome-derived myosin-like protein 1; GDMLP-1;
                                                                                                                                                         Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                skeletal
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l muscle
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2001WO-US00665.
2001WO-US00667.
2001WO-US00667.
2001WO-US00668.
2001WO-US00669.
2001WO-US00670.
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2001WO-US00663.
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2001WO-US00661.
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2000US-236359P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chromosome 22; gene therapy; vacci
disorder; amplicon; screening; ss.
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93.8%;
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                                                                                                                                                                                                                                                                       Hanzel
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Pred. No. 1.4e+02;
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                                                                                                                                                                                                                                                                       DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                       English.
                                                                                                                                                                                                                                                                       Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapy; vaccine; heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence SEQ
                                                                                                                                                                                                                                                                       Chen
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Matches 15
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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27-SEP-2000;
04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                     New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N.B. The sequence data for this patent did not specification, but was obtained in electronic
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                                                                                                                                                                                                                                                                                                                                                                                                                                    skeletal
                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart;
muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human GDMLP-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAY-2002 (first entry)
            myosin-like
                                                                                                                                                                  30-JAN-2001;
30-JAN-2001;
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                                                                     2002-179446/23.
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                                                                                             Ji Y,
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                                                                                                                     AEOMICA
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                                                                                                                                                                                                                                                                                                                                                                                                                                     muscle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard; DNA; 17
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2001WO-US00662.
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2001WO-US00665.
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2001WO-US00666.
2001WO-US00667.
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           protein
                                                                                                                                          2001WO-US00668.
2001WO-US00669.
2001WO-US00670.
2001US-266860P.
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                                                                                             Penn SG,
                                                                                                                     INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                  chromosome 22; gene disorder; amplicon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.0%;
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                                                                                            Hanzel
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Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO:5
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                                                                                             Rank
                                                                                                                                                                                                                                                                                                                                                                                                                                    screening; ss
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                                                                                             Chen
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                                                                                             Shannon
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Disclosure;

SEQ

ij 9002;

214pp;

English

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RESULT 122
ABZ61957/c
ID ABZ619
XX ABZ619
XX ABZ619
XX Human
XX Human
XX Human
XX Homo E
PN WO2002
YX WO2002
YX HOMO E
YX ADZ9-MAN
PR 29-MAN
PR 10-SEI
XX PR 10-SEI
XX WPI;
XX WPI;
XX WPI;
XX WPI;
XX WPI;
XX HER2,
XX HER2,
XX HER2,
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CC substrates, to provide initial substrates for the recombinant engineering
CC of hGDMLP-1 protein variants having desired phenotypic improvements, and
CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may
CC be used as immunogens to raise antibodies that specifically recognise
CC hGDMLP-1 proteins, as standards in assays used to determine the
CC concentration and/or amount specifically of hGDMLP proteins, as specific
CC deficiency in hGDMLP-1 production, and in vaccines or for replacement
CC deficiency in hGDMLP-1 production, and in vaccines or for replacement
CC therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for
CC diagnosing a disorder associated with the expression of hGDMLP-1, in
CC particular heart and skeletal muscle disorders. hGDMLP-1 is localised to
CC chromosome 22. The present sequence represents an oligomer used in the
CC screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; ribozyme; short interfering RN/enzymatic nucleic acid; H-Ras; N-Ras; anti-rheumatic; cancer; AIDS; 88.
                                      Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encod
                                                                                                                                                                                                             29-MAY-2001; 2001US-294140P.
06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
                                                                                                                                                                                                                                                                                                                                  05-DEC-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human H-Ras DNAzyme target #748.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABZ61957;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic can be used as probes to detect, characterise
                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                            29-MAY-2002; 2002WO-US16840
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABZ61957 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence.
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                                                                                                2003-140484/13
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                Ger, modulates the expression of a nucleic acid encoding H-Ras, N-Ras, and human deficiency virus sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   interfering RNA; siRNA; HER2; K
; H-Ras; N-Ras; HIV; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.4; DB
Pred. No. 1.4e+(
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   K-Ras;
c; anti-HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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RESULT 123
AAZ41119
ID AAZ411
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OS AAPI
XX AAZ41
XX AAZ4
XX AAZ41

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cowsert |
Ohasi C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ra human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66524, ABZ66520 - ABZ66524, ABZ66524 - ABZ66524, ABZ66524, ABZ66525 - ABZ66524, ABZ66524 - ABZ66524, ABZ66524 - ABZ66524, ABZ66524 - A
                                                                                                                                                                                                                                                                                                                                                            Identifying compounds which modulate expression of nucleic acids, un
to provide compounds having defined physical, chemical or bioactive
properties, e.g. antisense activity -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                 Example 27; Page 108; 264pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-620446/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-APR-1998;
28-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      target validation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identification; genetic target; gene modulation; human; probe; antisense oligonucleotide; phosphorothioate; PCR primer; nucleotide sequence-based technology; antisense drug discovery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human G-alpha-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequences for the human ribozymes of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1722 GGGCAAGCCCCTGGGA 1737
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C, Wyatt JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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98US-0067638.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BF, McNeil J,
, Borchers AH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ر:
و ر:
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Pred. No. 1.4e
0; Mismatches
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Vickers TA;
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A method has been developed of defining a set of compounds that modulat the expression of a target nucleic acid (tNA) sequence via binding of the compounds with the tNA sequence. The method comprises generating a library of virtual compounds in silico according to defined criteria, and evaluating in silico the binding of the virtual compounds with the tNA according to defined criteria. Also described are: (1) a method of defining a set of oligonucleotides (ONs) that modulate the expression of the virtual compounds.

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RESULT 124
ARAZ194
XX
XX
XX
ARZ194
AC ARZ194
XX
AC ARZ194
XX
DT 15-NOV
XX
Human;
KW Human;
KW phosph
XX
VS Synthe
OS Synthe
OS Synthe
OS Homo s
XX
US5951
XX
14-SEF
XX
14-SEF
XX
14-DEC
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04-DEC
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PT Inhibi
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                                                                                                                                                                             nucleotides, targeted to a nucleic acid molecule encoding human G-alpha-11. AAZ19468 to AAZ19547 resent human G-alpha-11 phosphorothioate antisense oligonucleotides given in the present invention. The oligonucleotides may be useful for the treatment of
                                                                                                                                                                                                                                                                                                                                                                                     Inhibitory antisense compounds useful for associated with G-alpha-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               exemplification of the present invention.
                                                                                                     Sequence 18
                                                                                                                                                        diseases associated
                                                                                                                                                                                                                                                                                     The present invention
                                                                                                                                                                                                                                                                                                                                        Example 15;
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                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGGCCGTGAAGATGTT 1829
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                                                                                                                                                                                                                                                                                                                                     Column 40; 38pp; English.
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                                                                                                     BP; 3
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  Conservative
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                        1.0%;
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                                                                                                     G; 6 T; 0 other;
Score 14.4; DB 1;
Pred. No. 1.5e+02;
0; Mismatches 1;
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Pred. No. 1.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                   the treatment
                                                Length 18;
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Gaps
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RESULT 125
ABK49070
ID ABK4907
XX ABK490
XX Cofact
XX Cofact
XX Casset
XX Synthe
XX Synthe
XX W02002
XX W11-SEI
XX H1-SEI
XX H1-SEI
XX WPI;
PT Cofact
PT in one
PT Cofact
PT in one
PT Cofact
PT Site a
XX (RUTF
XX WPI;
PT Cofact
PT in one
CC 2,5-d:
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CC acid a
CC acid
                      The present invention relates to a new method of producing mutant CC 2,5-diketo-D-gluconic acid reductase enzymes with altered cofactor CC dependency. The method of the invention involves identifying a cofactor CC specificity site in a wild type 2,5-diketo-D-gluconic acid reductase CC enzyme and mutating an amino acid in the identified cofactor specificity site so that cofactor dependency of reactions catalysed by the enzyme is CC altered. The method is useful for producing mutant 2,5-diketo-D-gluconic CC acid reductase enzymes with altered cofactor dependency and as such is CC useful in metabolically engineered organisms to produce 2-keto-L-gluconic CC acid from glucose in a single fermentation step. The mutant 2,5-diketo-D-gluconic acid reductase enzyme produces 2-keto-L-gluconic acid from CC glucose in a single fermentation step. The mutant 2,5-diketo-D-glucose in a single fermentation step. The mutant 2,5-diketo-D-glucose in a single fermentation step in metabolically engineered corganisms. Flexibility in catalysing the enzymetic reaction by the enzyme CC dinucleotide) or NADDH provides advantages. Since the cost of NADPH is an order of magnitude greater than that of NADH, use of the enzyme of the CC invention in any in vitro system where cofactor must be purchased and CC provided for the enzymes provides a significant cost saving advantage. The mutant 2,5-diketo-D-gluconic acid reductase enzyme exhibits increased levels of apparent expression which leads to an increased rate of classification of 2-keto-L-gluconic acid The present mucleic acid segmence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Making mutant 2,5-diketo-D-gluconic acid reductase enzymes with altered cofactor dependency, for producing 2-keto-L-gluconic acid from glucose in one fermentation step, comprises mutating a cofactor specificity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2,5-diketo-D-gluconic acid reductase A enzyme; cofactor cofactor dependency; metabolically engineered organism; 2-keto-L-gluconic acid; glucose; single fermentation stecassette mutagenesis; restriction site; pATP003.xb; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-351864/38.
P-PSDB; AAU79911.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 34; 45pp; English
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/product= "Wild-type pATP003.xb
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Best Local S
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Best Local (
                                                                                                                            A method for treating airway disease in a subject has been produced, which involves the topical administration of an essentially adenosing free antisense oligonucleotide (ON) to the airway epithelium of the subject. The present sequence is an antisense oligonucleotide specific for the human IL6 receptor. The method can be used to treat airway diseases such as cystic fibrosis, asthma, chronic obstructive pulmonary disease, bronchitis and other airway diseases characterised by an inflammatory response. By eliminating adenosine is the antisense ON, its liberation upon antisense degradation is prevented, thereby preventing adenosine-induced bronchoconstriction:
                                                                                                                                                                                                                                                                                                         Treatment of airway diseases such as asthma - by topically applying adenosine-free antisense oligo:nucleotide to airway epithelium of
                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                       Metzger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               represent the wild-type pATP003.xb restriction site #2 that encodes the wild-type pATP003.xb restriction site peptide #2. This sequence was used in the methods of the invention for cassette mutagenesis.
                                                                                                                       patients
                                                                                                                                                                                                                                                                       Example 5; Page 32; 71pp; English
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                                                                                                                                                                                                                                                                                                                                                                                               (UYEC-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                        07-JUN-1995;
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                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             airway epithelium; adenosine free; cystic fibrosis; obstructive pulmonary disease; bronchitis; interleukin; ss.
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GAGCCCGGAGGCCA 1
                    GAGCCCGGAGGCCA 2016
                                                                                               15
                                                                                                                      with hyper-reactive airways.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   receptor antisense oligonucleotide
                                                                                               BP;
                                                Conservative
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Pred. No. 1.5e+
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                                                Mismatches
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                                            1.4e+02;
hes 0;
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GAGCCCGGAGGCCA 2016 GAGCCCGGAGGCCA 1

14

Matches Query Match

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Conservative

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Mismatches

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Gaps

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Local

Similarity

100.0%;

Score 14; Pred. No.

1.4e+02 DB 1;

Length 15 Indels

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AAX54048/c
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                                                                                                                                              antisense treatment of diseases and conditions. Typical diseases and conditions are those associated with impaired respiration and inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, acute asthma, allergies, asthma, imprespiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, pulmonary vasoconstriction, emphysema, chronical pulmonary hypertension, pulmonary vasoconstriction.
                                                                                                                                                                                                                                                                           non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides may be derived from sequences AAX55272-74. These multiple target oligonucleotides (specifically AAX55180-271) can be used for the
                                     obstructive pulmonary disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g. colon cancer, breast cancer, lung cancer, panoreatic cancer, hepatocellular carcinoma, kidney cancer, melanoma hepatic metastases, as well as all types of cancers which may metastor or have metastasized to the lungs, including breast and prostate can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX54048 standard; DNA;
    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasoconstriction; emphysema;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotide; multiple target; antisense treatm impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                   The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 50; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense oligonucleotides used in treatment of, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-229400/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-JUN-1998;
17-SEP-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   colon cancer; breast cancer; lung cancer; pancreatic cancer;
hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human IL-6 receptor antisense oligonucleotide fragment
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    BP;
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0 A;
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97US-0059160.
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    3 T; 0 other;
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                                                                                                                                                     emphysema, chronic
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                                                                                         melanoma,
                                                                metastasize
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RESULT 128
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                     CC (I) can have respiratory, bronchodilator, antiinflammatory, analysesic, cc mmunosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC The antisense oligonucleotides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with cc lung/respiratory disorders and malignancies, such as stimulating and cactivating peptide factors and transmitters, transcription factors, immunoglobulins and antibodies, antibody receptors, cytokines and cc immunoglobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, binding proteins, adhesion molecules and their receptors, cytokine and cc receptors, binding proteins and non-nervous system errous system (CNS) and peripheral nervous and non-nervous system peptide transmitters, defensins, growth factors, vasoactive peptides and creceptors, binding proteins and malignancy associated proteins. The creceptors binding proteins and malignancy associated proteins. The creceptors binding proteins and malignancy associated proteins. The creceptors binding proteins and malignancy associated proteins. The cand/or bronchoconstriction) and/or lung inflammation, allergy(ies) and/or surfactant hypoproduction which are associated with a disease or condition selected from pulmonary vasoconstriction, inflammation, callergies, asthma, impeded respiration, respiratory distress syndrome (C and/or cancer, AAFJAGA14 to AAFJAGA1 represent human polyvuiclerief identificated in and variable to the condition and variable to condition constriction constriction beloated to condition polyvuiclerief identification condition and condition and condition condit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions - \frac{1}{2}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes low adenosine (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pulmonary hypertension; emphysema; pulmonary transplantation rejection; chronic obstructive pulmonary disease; pulmonary infection; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 14; Page 209; 1592pp; English.
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                                                                                               The present invention describes a new composition comprising an CC antisense oligonucleotide (ON) with low adenosine (up to 15%), which CC targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antiallergie, and/or can insthmatic, cytostatic and analyssic activities. The compositions are useful for the treatment of diseases associated with inflammation, compared airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating ce.g. ischaemic conditions, pulmonary vasoconstriction, allergies, casthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, content of the seast and prostate cancer. The reduction of the adenosine content of the lungs, including the seast and prostate cancer. The reduction of the adenosine content of the lungs in the later of the lungs in the later of the lungs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches 14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15 BP; 0 A; 7 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA33492
                                       the ONs reduces side effects. The A-containing ONs break down with the release of deoxyadenosine which activates adenosine receptors causing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Low adenosine antisense oligonucleotide SEQ ID
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Pred. No.
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Best Local
                                                                                                                                                                                                                               haemopoietin receptor protein family NR8 genes. The NR8 family sequences were initially searched for comparison on a nucleic acid database with the nucleic acid probe sequence TGGAGYNNNTGGAGY encoding the amino acid sequence Trp-Ser-Xaa-Trp-Ser. The sequences ARZ59258-Z59300 and ARZ90816-Z99925 represent specific examples of probe sequences used in the search. Antibodies to the NR8 family proteins are used for the diagnosis of blood formation disorders. Compounds identified as binding to the proteins are used for the treatment of such disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 sequences are also called SEQ ID NO:1 to 2815, and then the last differ from the previously named sequences. SEQ ID NO:11 to 1680 (AAA32323 to AAA33992) are specifically claimed ONs from the present invention. N.B. Sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequences given in the sequence listing.
                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hemopoietin receptor protein family NR8 formation disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to the isolation of sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 41; 176pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CHUG-) CHUGAI RES INST MOLECULAR MEDICINE INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15 BP; 0 A; 7 C; 5 G; 3 T; 0 other;
1581 CTCCATGAACTCCA 1594
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                                                                                                           Similarity
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                                                                               Conservative (
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98JP-0297409.
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k; Pred. No. 1.4
0; Mismatches
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hes 0;
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                                                                                                                                    DB 1;
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RESULT 131
AAF52667
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                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Pactor 1 receptor; IGF-1; pityriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF52667;
                                                                                                                                                                                       sclerotic disease, kidney disease, hyperproliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 8; Page 84; 201pp; English
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                                                                                    Local
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                                                                                  Similarity
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AAH95775
AC X I
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AMF45151 and AMF45153-F45161). The method is useful for ameliorating the effects of AMF45153-F45161) is pityriasis, ruba, pilaris, serbornhoea, keloids, particular and provided that the context of the present invention (see AMF45151).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFB-2; IGFBB3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell
AAH95775
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 0 A; 4 C; 6 G; 5 T; 0 other;
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                                                          AAH95775 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sclerotic disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to a method for ameliorating the effects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 8; Page 84; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             proliferation and/or inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MURD-) MURDOCH CHILDRENS RES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IGF-I oligonucleotide #3631.
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                                                                                                                                                                                                                                                                                            2332 TEGTCCTTCEGET 2345
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    kidney disease, hyperproliferation of the inside of
any other hyperplasia.

                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
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100.0%; Pred. No.
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AAH95776
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides nucleic acid molecules capable of downregulating the expression of the human checkpoint kinase-1 (Chk1) gene. These may be antisense or ribozyme sequences, and are useful in the treatment of diseases associated with conditions affected by Chk1 levels, including cancer. The present sequence is an oligonucleotide described in
                                                                                                                                                                                           09-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 6 A; 3 C; 6 G; 2 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel nucleic acid molecule e.g., ribozymes or antisense nucleic acid molecules, which downregulates expression of a checkpoint kinase-1 gene, useful for treating colorectal, lung, breast or prostate cancer
                                                                                                                                                                                                                   AAH95776;
                                                                                                                                                                                                                                                                                                                                                                                                                      the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-FEB-2000; 2000US-0179983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-FEB-2001; 2001WO-US03504.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human Chkl ribozyme substrate SEQ ID NO: 1200.
                                  02-FEB-2001; 2001WO-US03504.
                                                          09-AUG-2001
                                                                                  WO200157206-A2
                                                                                                         Homo sapiens.
                                                                                                                                RNA cleavage;
                                                                                                                                            Human; checkpoint kinase-1; Chkl; antisense; ribozyme; gene therapy;
                                                                                                                                                                  Human Chk1 ribozyme substrate SEQ ID NO: 1201
                                                                                                                                                                                                                                         AAH95776 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 88; 115pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cleavage;
                                                                                                                                                                                                                                                                                                                          1730 CCCTGGGAGAAGGT 1743
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12; Conserv
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                                                                                                                                                                                          (first entry)
                                                                                                                                 cancer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                       invention.
                                                                                                                                                                                                                                                                                                                                               Score 14; DB 1; Le
Pred. No. 1.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                        Length 17;
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03-FEB-2000; 2000US-0179983

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RESULT 135
AAH95777
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Best Local :
         downregulating the expression of the human checkpoi
gene. These may be antisense or ribozyme sequences,
                                                                                      Novel nucleic acid molecule e.g., ribozymes or antisense nucleic acid molecules, which downregulates expression of a checkpoint kinase-1 gene, useful for treating colorectal, lung, breast or prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention provides nucleic acid molecules capable of downregulating the expression of the human checkpoint kinase-1 (Chk1) gene. These may be antisense or ribozyme sequences, and are useful in the treatment of diseases associated with conditions affected by Chk1 levels, including cancer. The present sequence is an oligonucleotide described in
                                                        Claim 4;
                                                                                                                                                             Fattaey
                                                                                                                                                                                                                                              02-FEB-2001; 2001WO-US03504
                                                                                                                                                                                                                                                                       09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                    Human; checkpoint kinase-1; Chkl; antisense; ribozyme; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                              Human Chkl ribozyme
                                                                                                                                                                                                                                                                                                                                                                                                     09-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RIBO-)
                                                                                                                                                                                                 (RIBO-)
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                                                                                                                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                                                                           RNA
                                                                                                                                                                                                                                                                                                                                                                                                                            AAH95777;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             molecules,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel nucleic acid molecule e.g.,
                   present invention provides nucleic acid molecules capable or nregulating the expression of the human checkpoint kinase-1
                                                                                                                                                                                                                                                                                                                                          cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1730 CCCTGGGAGAAGGT 1743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              luding cancer. The present sequence exemplification of the invention.
                                                                                                                                         2001-496922/54
                                                                                                                                                                                                                                                                                                                   sapiens
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                                                                                                                                                                                     RIBOZYME PHARM FATTAEY A R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ₽,
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                                                                                                                                                               Ŗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                     Page 88; 115pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCCUGGGAGAAGGU 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 5 A; 3 C; 7 G; 2 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                               Jarvis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nich downregulates
for treating color
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                                                                                                                                                                                                                                                                                                                                          cancer;
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                                                                                                                                                                                                                                                                                                                                                                              substrate
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85.7%;
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                                                       English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             le e.g., ribozymes or antisense nucleic acid lates expression of a checkpoint kinase-1 colorectal, lung, breast or prostate cancers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14;
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                                                                                                                                                                                                                                                                                                                                                                            SEQ
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                                                                                                                                                             Booher RN,
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gene.

of diseases associated

with

conditions affected

and are useful

(Chk1)

by Chk1 levels,

for expressing

protein variants having desire ing the proteins. The hGDMLP-1

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Matches 12; Conserv
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          The present invention describes a human genome-derived myosin-like protein 1 (hgDMLP-1). The protein and polymuces of hgDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantity hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and
                                                                                                                                                                                                                                                                                                                                           30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                        New polypeptide, for raising antibodies that proteins, or as specific biomolecule capture surface-enhanced laser desorption ionization, myosin-like protein hGDMLP-1
                                                                                                                                                                                                                                                                                                                                                                                                   21-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; genome-derived myosin-like protein 1; GDMLP-1;
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                                                                                                                                                                                                                             (AEOM-)
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30-JAN-2001;
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                                                                                                                                                                                                                            AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCCUGGGAGAAGGU 14
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                                                                                                   SEQ ID 6888; 214pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                             2001WO-US00669.
2001WO-US00670.
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                                                                                                                                                                                                       Hanzel
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ
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                                                                                                                                                                                                       DR,
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                                                                                                                                   recognize hGDMLP-1 probes for comprises human
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heart disease;
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                                                                                                                                                                                                       Shannon ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ID NO:6888
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RESULT 137
ABN06897/c
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Matches
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21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 protesins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP protesins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N.B. The sequence data for this patent did not specification, but was obtained in electronic at ftp.wipo.int/pub/published_pct_sequence.
                                                                                                                                                                                                                                                                                                                                                                            25-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                           06-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6889.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABN06897;
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                                           2002-179446/23.
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                                                                                                       AEOMICA
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2001WO-US00666.
2001WO-US00667.
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                                                                                                                                 2001US-266860P
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2001WO-US00670
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2001WO-US00661.
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2000US-234687P.
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                                                                      Penn SG,
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100.0%; Pred. No.
                                                                         Hanzel DK,
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hes 0;
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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for

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RESULT 138
ABN06898/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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           30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                      Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N.B. The sequence data for this patent did not specification, but was obtained in electronic f at ftp.wipo.int/pub/published_pct_sequence.
                                                                       30-JAN-2001;
30-JAN-2001;
                                                                                                    04-OCT-2000;
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                                                                                                                    21-SEP-2000;
27-SEP-2000;
                                                                                                                                                  26-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                    Human GDMLP-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein 1 (hGDMLP-1). The protein and polynucrevities sequents hGDMLP-1 can be used in gene therapy and vaccine production.
                                                                                                                                                                                  25-MAY-2001; 2001WO-US16981.
                                                                                                                                                                                                                                            WO200192524-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          surface-enhanced laser desorption ionization, comprises human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention
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           2000GB-0024263.
2001WO-US00661.
2001WO-US00662.
2001WO-US00663.
2001WO-US00665.
2001WO-US00665.
2001WO-US00666.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 1 A; 6 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID 6889; 214pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein hGDMLP-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention
                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                     2000US-236359P
                                                                                                                                                  2000US-207456P
                                                                                                                                                                                                                                                                                                                                                                scanning SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The protein and polynucleotide sequences of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 describes a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human genome-derived myosin-like
                                                                                                                                                                                                                                                                                                                                                                  NO:5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; L
                                                                                                                                                                                                                                                                                                                                                                  sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      form part of the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO:6890.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         directly from WIPO
                                                                                                                                                                                                                                                                                                                       hGDMLP-1; heart; heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
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RESULT 139
ABN06899/c
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30-JAN-2001;
30-JAN-2001;
05-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ည
                                                                                                                                          Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart,
                                                                                                                                                                                                                29-MAY-2002
                                                                                                                                                                                                                                                                              ABN06899 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; SEQ ID 6890; 214pp;
                  06-DEC-2001
                                                  WO200192524-A2
                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                ABN06899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        myosin-like protein hGDMLP-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-179446/23.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention
                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                            1358 CGCCTGGAAGAGAA 1371
                                                                                                                                                                                                                                                                                                                                                              15
                                                                                                                                                                                                                                                                                                                                                                                                                              14;
                                                                                                                 myosin;
l muscle
                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                              CGCCTGGAAGAGAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 1 A; 6 C; 5 G; 5. T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001WO-US00669.
2001WO-US00670.
2001US-266860P.
                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001WO-US00668.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Penn SG,
                                                                                                                                                                              17-mer scanning
                                                                                                                 chromosome 22; gene disorder; amplicon;
                                                                                                                                                                                                                                                                                                                                                                                                                                             1.0%; Score 14;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hanzel
                                                                                                                                                                                                                                                                              ВP
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                                                                                                                                                                              SEQ ID NO:5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rank DR,
                                                                                                                              therapy; vaccine; heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                             1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                              sequence SEQ ID NO:6891.
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                                                                                                                                                                                                                                                                                                                                                                                                                            <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
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RESULT 140
AAQ24060
ID AAQ240
XX
AC AAQ240
XX
DT 08-JUN
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08-JUN-1992 AAQ24060; AAQ24060

(first entry)

standard; RNA; 17

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Best Local S
Matches 14
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21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                         The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption instation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement
                                                                                                                                                                                                              particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1
                                                                                                                      Sequence
                                                                                                                                                  at ftp.wipo.int/pub/published_pct_sequence.
                                                                                                                                                                  N.B. The sequence data for this patent did not specification, but was obtained in electronic f
                                                                                                                                                                                                                                                            therapy. The polynucleotide sequences encoding hGDMLP-1 may be usudiagnosing a disorder associated with the expression of hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gu Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure;
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30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAY-2001;
                                                                                                                                                                                                 invention.
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30-JAN-2001;
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                           1358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2002-179446/23.
14
                                                        l Similarity
14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AEOMICA INC
                 CGCCTGGAAGAGAA 1371
CGCCTGGAAGAGAA 1
                                                                                                                        17
                                                        Conservative (
                                                                                                                     BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID 6891; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001WO-US00665.
2001WO-US00666.
2001WO-US00667.
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2001WO-US00663.
2001WO-US00664.
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                                                                                                                     6 C; 4 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hanzel DK,
                                                          <u>,</u>
                                                                       Score 14; DB 1;
Pred. No. 1.6e+
                                                                                                                     6 T;
                                                          Mismatches
                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rank DR,
                                                                                                                      other;
                                                        1.6e+02;
                                                                                                                                                                    format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chen
                                                                                                                                                                                   form
                                                                                      Length 17;
                                                          Indels
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                                                                                                                                                                  part of the printed directly from WIPO
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                                                                                                                                                                                                                                                                              be used
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RESULT 141
AAX74934
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Best Local Similarity
Matches 13; Conser
                                                                                                                                                                                                                                                                                                                                                                                                         structures are in the loop region and the U-rich bubble in the upper stem. This 17-mer was synthesised and can hybridise to a 14-mer (see AAQ24061) to mimic the known HIV-1 tat recognition sequence but without the apical loop. In an assay, the 17-mer plus 14-mer structure competed satisfactorily with full-length (59-mer) TAR for binding to tat. See AAQ21427-Q21435 for TAR mutants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Key
misc_structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New HIV growth inhibiting oligo:nucleotide(s) - comprising binding sequences capable of binding to tat protein within and in assays to identify cpds. with tat binding
                                                                        Vascular endothelial growth factor receptor; YEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The HIV-1 TAR stem-loop sequence (see AAQ21425) was compared to that from HIV-2 (see AAQ21426). The only regions common to the two TAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Fig 18c; 89pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Karn
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        trans-activation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human immunodeficiency virus; tat protein; AIDS; hairpin loop;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Artificial HIV-1 TAR sequence
WO9715662-A2.
                                                               fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                Mouse flt-1
                                                                                                                                                         28-JUL-1999
                                                                                                                                                                                                            AAX74934 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17
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                                                                                                                                                                                                                                                                                                  1490 AGCCAGACTTCAGCAGC 1506
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                                                    liver
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                                                                                                                                                                                                                                                                            AGCCAGAUUUGAGCAGC
                                                                                                                                                                                                                                                                                                                                                                                  BP; 5
                                                                                                                                VEGF
                                                                                                                                                                                                                                                                                                                                Conservative
                                                 kinase 1; ss:
                                                                                                                                                         (first entry)
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                                                                                                                               receptor hammerhead ribozyme substrate #462
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      responsive
                                                                                                                                                                                                                                                                                                                                                                                  A; 4 C; 5 G; 3 U; 0 other;
                                                                                                                                                                                                                                                                                                                                         1.0%;
76.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "U-rich bubble. Base pairs to nucleotides 6-10 of AAQ24061"
                                                                                                                                                                                                                                                                              17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The only regions common to the two TAR
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                                                                                                                                                                                                                                                                                                                               Score 13.8; DB 1;
Pred. No. 1.8e+02;
2; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      region; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          containing U-rich bubble
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dingwall C;
                                                                                                                                                                                                                                                                                                                                                        Length 17;
                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA
cells,
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                                                                                                                                                                                                                                                                                                                              Gaps
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11-JAN-1996; 26-OCT-1995;

96US-0584040. 95US-0005974. 96WO-US17480.

(CHIR) CHIRON CORP.
(RIBO-) RIBOZYME PHARM INC

25-OCT-1996; 01-MAY-1997.

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vescular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 2 A; 4 C; 4 G; 7 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-JAN-1996;
26-OCT-1995;
                                                                                                                                                                                                                           Vascular endothelial growth factor receptor, VEGF receptor; flt-1; flk-1; KDR; hammerhad ribozyme; hairpin ribozyme; cleavage; tlk-1; KDR; hammerhad ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                    Mouse flt-1 VEGF receptor hammerhead ribozyme substrate #463.
                                                                                                                                                                                                                                                                                                                                                                                          AAX74935 standard; RNA;
                                                                                                                                                         WO9715662-A2
                                                                                                                                                                                    Mus sp.
                                                                                                                                                                                                                                                                                                                                 28-JUL-1999
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(RIBO-) RIBOZYME PHARM INC.
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                                                                                                         Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase i; kinase insert domain containing receptor;
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                                                    Claim 4; Page 169; 218pp; English
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present invention

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acid molecules which modulate

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angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.

expression ic examples

The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour

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Best Local :
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                                                                                                                                The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psociasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1, KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-1ke tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
                                                                                                                                                                                                                                                                                                               Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
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26-OCT-1995;
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                                                                                                        Sequence 17
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                        2279 GGATGGCTCCAGAAGCC 2295
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RIBOZYME PHARM INC.
                                                                 Similarity
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95US-0005974.
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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RESULT 146

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mRNA stability - useful for treating e.g. tumour angioge
psoriasis, rheumatoid arthritis, etc., in a human patient
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26-OCT-1995;
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Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage;
                                                                                                                                                                                                                                                                            AAX73029 standard; RNA; 17 BP
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(RIBO-) RIBOZYME PHARM INC.
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95US-0005974.
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                                                                                             receptor hammerhead ribozyme substrate #462
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Pred. No. 1.8e+02;
7; Mismatches 2;
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RESULT 148
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Best Local
                                                                                                                               Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-1ike tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
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   25-OCT-1996;
                               01-MAY-1997.
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                                                            WO9715662-A2
                                                                                                                                                                                                        Mouse flk-1 VEGF receptor hammerhead ribozyme substrate #463
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26-OCT-1995;
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Similarity 58.8%;
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95US-0005974.
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Pred. No. 1.8e+02;
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RESULT 14
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                                                                                                              11-JAN-1996;
26-OCT-1995;
                                                                                                                                                                                                                                                                                 Vascular endothelial growth factor receptor; VBGF receptor; flt-1; flk-1; KTR; hammerhad ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
              WPI; 1997-259017/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
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26-OCT-1995;
                                                                  (CHIR ) CHIRON CORP.
(RIBO-) RIBOZYME PHARM INC
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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing
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                       The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
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AAX71455
                            AAX71455 standard; RNA; 17
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                                                                                                                                                                                                             BP; 3 A; 3 C; 5 G; 6 U; 0 other;
                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    McSwiggen J, Pavco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          96US-0584040.
95US-0005974.
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                                                                                                                                                                  1.0%;
                                                                                                  17
                            ВP
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د
                                                                                                                                                      Score 13.8; DB 1;
Pred. No. 1.8e+02;
5; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P, Stinchcomb
                                                                                                                                                       2;
                                                                                                                                                                                 Length 17;
                                                                                                                                                      Indels
                                                                                                                                                      0
                                                                                                                                                       Gaps
                                                                                                                                                       0
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AAA18893
ID AAA188
XX AAA188
AC AAA188
XX ID 19-JUN
XX IP-JUN
XX IP-JUN
XX INTEGER
XX INTEGER
XX INTEGER
XX INTEGER
XX AAA188

Human TIE-2

substrate sequence SEQ ID NO:2119.

19-JUN-2000

(first entry)

AAA18893 standard;

RNA;

17

Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis;

macular degeneration; inflammation; neovascular glaucoma;

age related

밁 Ś

Matches

10;

<u>ა</u>

Mismatches

0

Gaps

0

2112 CATGGAGTACTTGGCTT 2128

CAUGGAGUUCUUGGCAU

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Query Match
Best Local
                                                                                            The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                       Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                      (CHIR )
                                                                Sequence 17 BP; 3 A; 3 C; 5 G; 6 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-JAN-1996;
26-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vascular endothelial growth factor receptor; VBGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                           Claim 4; Page 111; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                      Escobedo J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W09715662-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               foetal liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KDR VEGF receptor hammerhead ribozyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                      CHIRON CORP.
RIBOZYME PHARM INC.
                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                      McSwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96US-0584040.
95US-0005974.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96WO-US17480.
                  1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                      Pavco P,
                   Score 13.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                        Stinchcomb
8; L.
1.8e+02;
2;
                                Length 17;
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RESULT 155
AAZ00939/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA23422 represent their corresponding target sequences. The ribozymes the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT; integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2, integrin subunit alpha-6, or integrin subunit beta-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA1675 to AAA1761 to AAA1762 represent ribozyme sequences for RNT, and AAA17168 to AAA17631 to AAA17623 represent tribozyme sequences for RNT, and AAA17168 to AAA17634 represent tribozyme sequences for Tie-2, and AAA19087 to AAA19154 represent ribozyme sequences for Tie-2, and AAA19086 and AAA19155 to AAA20361 and AAA19015 to AAA21506 to AAA21501 to AAA21505 represent ribozyme sequences for Tie-2 and AAA19086 and AAA19223 to AAA21506 and AAA21501 to AAA21505 represent ribozyme sequences for Tie-3 and AAA21508 to AAA21508 to AAA21508 represent their corresponding target sequences; AAA21506 to AAA21689 represent their corresponding target sequences; AAA21506 to AAA21687 and AAA23263 to AAA23342 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA2362, AAA23343 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-OCT-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
PG1 gene; biallelic marker; PCR primer; PG1-related biallelic marker;
                                         PCR primer
                                                                                  27-SEP-1999
                                                                                                                           AAZ00939
                                                                                                                                                                AAZ00939 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 4 A; 1 C; 6 G; 6 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 56; Page 123; 305pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pavco PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-MAR-1999;
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                                                                                                                                                                                                                                                                                                              2344 GTGTTAATGTGGGAGAT 2360
                                                                                                                                                                                                                                                                          Н
                                                                                                                                                                                                                                                                                                                                                          Similarity
9; Conserv
                                                                                                                                                                                                                                                                          GUGUUACUAUGGGAGAU 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Roberts E,
                                         moPGrace5R444 for PG1
                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98US-0079678
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                                                                                                                                                                                                                                                                                                                                                                           1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jarvis T,
                                                                                                                                                                                                                                                                                                                                                       Score 13.8; DB 1; Length 17; Pred. No. 1.8e+02; 6; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Coeshott C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         McSwiggen JA;
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                                                                                                                                                                                                                                                                                                                                                     Gaps
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27-MAY-1999 WO9925815-A2

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RESULT 156
AAX76853/c
ID AAX768
XX
AC AAX768
XX
AC AAX768
XX
DT 05-AUG
DX
DC PCR pr
XX
VCOULTA
KW COULTA
KW PCR pr
XX
EV PCR PCR
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WO PCR PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a mammalian PG1 gene and protein, and a set of CPG1 biallelic markers. The PG1 polynucleotide and biallelic markers are used in a hybridisation assay, a sequencing assay or in an CPG1 callele-specific amplification assay for determining the identity of a conclectide at a PG1-related biallelic marker. The methods can be used to detect and to assess the risk of developing cancer or prostate specific antigen (PSA) dosage. However, the effectiveness of this is limited due to its inability to discriminate between malignant and non-malignant affections of the organ. A need exists for both a reliable diagnostic procedure which would enable early-stage diagnossis, and for preventative and curative treatments of the disease. The PG1 gene can be used for detection of prostate cancer, and the risk of developing it in the future, and can also be used to determine therapies for the disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                      Transcription unit; MARK2 kinase; rsk3 kinase; regulatory region; T66Bk; contraceptive; Responder/Distorter signalling cascade; t-Responder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX76853 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-405178/34.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-SEP-1998;
22-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9932644-A2
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                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                     PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer for cloning of T66Bk gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX76853;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       derived from it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Use of a prostate cancer associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUL-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2092 ACCTACCAGCTGGCCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17 ACCTACCTGCTGGCCTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%;
Similarity 88.2%;
15; Conservative
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                                                                                                                                                                                                                                                                                                                         88
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97US-0996306
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 13.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chumakov I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .8e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and biallelic markers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cohen D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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ARSSULT 157
AAV91368/C
ID AAV913
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AC AAV913
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AC AAV913
XX
DT 18-FEE
XX
DE Human
XX
Human

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence is a PCR primer used in the cloning of the T66Bk gene.

CC The invention related to a nucleic acid molecule (I) comprising a

CC transcription unit encoding in its 5' portion a kinase having a homology

CC to MARK2 kinase and the 3' portion of the nucleotide sequence has a high

CC containing (I) are useful for production of offspring. T66Bk, its

CC regulatory region, recombinant DNA, vectors, host cells, antibodies,

CC etc., are useful for the isolation of receptors on the surface of sperm

CC recognising attractants of the egg cell for the development and/or

CC production of contraceptives. They can also be used to identify chemicals

CC or biological compounds able to trigger the (premature) activation or

CC inhibition of the Responder/Distorter signalling cascade, or to identify

CC and isolate receptors and other members of the cascade that bind the

CC expression products. The methods for detecting the sperm of the

CC transgenic animal, and selecting against (I) also provide a means for

CC distorting the transmission ratio of genetic traits by altering genes of

CC the Responder/Distorter signal cascade other than the t-responder. They

CC also allow distortion, to a non-Mendelian ratio, of the transmission of a

CC effenting the expression distortion of sex, from male mammals to their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                    target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification; synthesis; deprotection; purification; cancer; inflammation; psoriasis; non-hepatic ascites; infection; genetic drift; restenosis; rheumatoid arthritis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          offspring by expressing during spermatogenesis/spermiogenesis a gene involved in sperm motility and/or fertilisation. The genes and proteins involved in the responder phenotype and Responder/Distorter signalling cascade, as well as the inventive methods are advantageous in breeding strategies by allowing for specific selection of genetic traits and in particular, of sex.
                                                                                                                                                                                                                                                                                                                                             Human;
                                                                                                                                                                                                                                                                                                                                                                                                         Human C-raf target site nucleotide position 2752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV91368;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV91368 standard; RNA; 17
   12-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 3 A; 7 C; 3 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-MAR-1998;
18-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2272 GTCAAGTGGATGGCTCC 2288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17 GTGAAGTGGATGGCACC 1
                                                                                                                                                                                                                                                                                                                                          c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7; Page 59; 117pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           acids involved in the Responder phenotype
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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97EP-0120190.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.8; DB
Pred. No. 1.8e+(
0; Mismatches
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AAV91369/c
ID AAV913
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AC AAV913
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AC AAV913
XX
DT 18-FEB
XX
DE Human
XX
KW Human;
KW Human;
KW Carget
KW deliye
KW gurifi

18-FEB-1999

(first entry)

AAV91369 standard; RNA;

17

0

Human C-raf target site nucleotide position

2753.

target; substrate; catalyst; modulation; expression; Raf gendelivery; screening; identification; synthesis; deprotection Human; c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin

psoriasis; non-hepatic

gene; ascites;

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                                                                                                                                         CC capable of modulating a process in a biological system. The method comprises: (a) introducing into the system a random library of nucleic comprises: (a) introducing into the system a random library of nucleic comprises: (a) introducing into the system a random library of nucleic comprises: (a) introducing into the system a random library of nucleic cord catalysts (NAC) having a substrate binding domain (SBD), comprising comprises activity and catalytic domain (CD); and (b) identifying NAC consists where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules cof at least part of the SBDs in such systems. Nucleic acid molecules cof at least part of the SBDs in such systems. Nucleic acid molecules call and to cleave target nucleic acid, particularly for treating colls and to cleave target nucleic acid, particularly for treating consists, non-hepatic ascites and infection. They may also be used to detert genetic drift and mutations in diseased cells and to determine constitution of the Raf gene, are used to treat cancer, restenosis, consortiasis or rheumatoid arthritis, or generally any condition associated with the level of c-raf. Introduction of sugar/phosphate modifications consorting that can be used in the method, specifically for capitesent NACs that can be used in the method, specifically for
                                                                   Matches
                                                                                    Query Match
Best Local
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Parry T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons
                                                                                                                                       Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 177; Page 153; 259pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-NOV-1997,
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                                                                                                                                                                         modulating the expression of a Raf gene.
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17
                                                                   15;
                                                                                    Similarity
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                               GCGCCTGGAAGAGAAAA 1373
GCTCCTGGAAGACAAAA 1
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                                                                                                                                       BP;
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, Kisich K, Matulic-Adamic J, McSwiggen JA;
                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97US-0056808.
97US-0061321.
97US-0061324.
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97US-0051718.
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97US-0046059
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                                                                                                                                       A,
                                                                                                                                       4 C; 4 G; 7 U;
                                                                                    1.0%;
88.2%;
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                                                                 Score 13.8; D
Pred. No. 1.8e
0; Mismatches
                                                                                                                                       0 other,
                                                                                1.8e+02;
                                                                                                    DB 1;
                                                                                                  Length 17;
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RESULT 159
AAF04325/c
ID AAF043
XX
AC AAF043
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DT 16-FEB
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                                                                                                                                                                                                                            Query Match
Best Local :
                                                                                                                                                                                                               Matches
 16-FEB-2001
                               AAF04325;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-JUN-1997;
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                                                               AAF04325 standard; DNA; 17
                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 177; Page 153; 259pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-009494/01.
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Parry T, Reynolds M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           infection; genetic drift; restenosis; rheumatoid arthritis; ss.
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                                                                                                                                                                             1356 АССССТССААСАСАА 1372
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15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    new catalytic nucleic acid that modulates selected
                                                                                                                                                                                                               Conservative
(first entry)
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97US-0061324.
97US-0064866.
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97US-0051718.
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                                                                                                                                                                                                                          1.0%;
                                                                                                                                                                                                                                                                           4 C; 4 G; 7 U; 0 other;
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                                                               BP.
                                                                                                                                                                                                           Score 13.8; DB 1;
Pred. No. 1.8e+02;
0; Mismatches 2;
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mic J, McSwiggen
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                                                                                                                                                                                                                                          Length 17;
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                                                                                                                                                                                                               Indels
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RESULT 160
AAF04773/c
ID AAF0477
XX AAF047
XX AAF047
XX AAF047
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XX Ribozy
KW Ribozy
KW interf
XX Homo s
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Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Enzymatic and antisense nucleic acid inhibition of repressor geuseful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Blatt L,
                                       12-APR-1999;
                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                        AAF04773 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-647423/62
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                (RIBO-) RIBOZYME PHARM INC
                                                                11-APR-2000; 2000WO-US09721.
                                                                                                                    WO200061729-A2
                                                                                                                                                                     Ribozyme; erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss.
                                                                                                                                                                                                            Hammerhead ribozyme substrate #2289.
                                                                                                                                                                                                                                      16-FEB-2001
                                                                                                                                                                                                                                                               AAF04773
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; Page 98; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-APR-2000; 2000WO-US09721.
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                                                                                          L9-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                         1.0%;
nilarity 88.2%;
Conservative
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                                       9905-0129390
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Pred. No. 1.8e+02;
0; Mismatches 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 other;
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RESULT 161
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The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, ERR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved the production of erythropoletin, granulocyte colony stimulating facts protein and interferon alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factors are the production of erythropoietin, granulocyte colony stimulating factors are the production of erythropoietin, granulocyte colony stimulating factors are the production of erythropoietin.
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                                                                                                                                                                                                                                                                                                                            useful for producing e.g. granulocyte colony stimulating protein, interferon alpha and erythropoietin -
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                                                                                                                                                                                                                                                                        Claim 54; Page 135; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic and antisense nucleic acid inhibition of repressor genes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-647423/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RIBO-)
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nilarity 88.2%;
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Matches Query Match Best Local

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RESULT 162
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Matches 15; Conserv
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                                                                                                                                                                                                        The invention relates to a method of determining if a compound binds to a target RNA by treating a test compound with a reporter (R) labelled with a donor or acceptor group and labelled target RNA, labelled with the complementary donor or acceptor group, and measuring the fluorescence from fluorescent groups associated with a compound:target RNA complex in presence of the test compound and comparing the result with a standard. The oligonucleotides AAZ59070-Z59071 anneal to form a double stranded oligonucleotides AAZ59070-Z59071 anneal to form a double stranded oligonucleotide containing the HIV-1 trans-activation regulatory region (TRR) to which the HIV-1 Tat protein binds. The complex is labelled with 6-carboxyfluorescein and is used as a target for the binding of a labelled ADP-1 protein. Detection of the complex is by fluorescence resonance energy transfer (FRET). The method is used to identify compounds that interfere with interaction between the target RNA and ligands or proteins.
Sequence 17
                                                                                         and ligands or proteins. Compounds that are identified are potentially useful for treating infections (viral, bacterial or fungal), cancer and autoimmune diseases. The compounds are preferably directed to the TAR and RRE regions of human immunodeficiency virus RNA and inhibit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antiviral; antibacterial; antifungal; anticancer; detection; TAR; fluorescence resonance energy transfer; tat; HIV-1; Rev response autoimmune disease; trans-activation regulatory region; ss.
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                                                               viral replication.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Examples; Page 31; 82pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying compounds that bind to target RNA, potentially useful reating infections, tumors and autoimmune diseases
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nilarity 88.2%;
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99GB-0004790.
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      4 C;
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                                                                                                                                                                                                                                                                                                                          The present invention provides nucleic acid molecules capable of downregulating the expression of the human checkpoint kinase-1 (Chk1) gene. These may be antisense or ribozyme sequences, and are useful in the treatment of diseases associated with conditions affected by Chk1 levels,
Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme
                                                    Human NOGO
                                                                             12-MAR-2002
                                                                                                      ABK00844
                                                                                                                            ABK00844 standard; RNA; 17
                                                                                                                                                                                                                                                                                         Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel nucleic acid molecule e.g., ribozymes or antisense nucleic acid molecules, which downregulates expression of a checkpoint kinase-1 gene, useful for treating colorectal, lung, breast or prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fattaey AR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIBO-)
(FATT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; checkpoint kinase-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Chk1 ribozyme substrate SEQ ID NO: 430
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                                                                                                                                                                                                                                                                                                                 luding cancer. The exemplification
                                                                                                                                                                                                                 2194 AAAATAGCAGACTTTGG 2210
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                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                          AAAAUCUCAGACUUUGG 17
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                                                   Inozyme
                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000US-0179983.
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                                                                                                                                                                                                                                                                                                                  invention.
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                                                                                                                                                                                                                                                    Score 13.8; DB 1;
Pred. No. 1.8e+02;
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  ribozyme
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The invention relates to a nucleic acid molecule which down regulates coppression of a CD20 gene and a nucleic acid molecule which down CC regulates expression of a neurite growth inhibitor gene (NOGO).

CC The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CC DNAzyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a CC DNAzyme) an Inozyme (aleaving RNA with a NGN triplet), a zinzyme (cleaving RNA with a NGN triplet), a zinzyme CC motif) pr an amberzyme (cleaving RNA with a NGN triplet), a zinzyme CC coleaver RNA of CD20 in the presence of a divalent cation that is CC preferably Mg.2^+. Furthermore, it may be contacted with a cell to reduce CC cD20 activity of the cell and treat a patient having a condition CC associated with the level of CD20. The treatment may further comprise the CC use of one or more therapies. In particular, the CD20 targetting CC lymphoma. low-grade or follicular NnL, lymphoma, leukaemia, B-cell lymphoma (NnL), bulky Clymphoma. low-grade or follicular non-Hodgkin's lymphoma (NnL), bulky CC lymphoma low-grade or follicular NnL, lymphocytic leukaemia, HIV (human cC immunocytoma (IMC), small B-cell lymphoma, limmune CC immunocytoma (IMC), small B-cell lymphoma, limmune (CC immunocytoma (IMC), small B-cell lymphocytic leukaemia, immune CC divalent cation that is preferably Mg.2^+. Furthermore, the nucleic acid contacted with a cell to reduce NOGO gene in the presence of a creatment may further comprise the use of one or more therapies.

CC treat a patient having a condition associated with the level of NOGO. The treatment having a condition associated with the level of CVA.

CC central nervous system (NOGO targetting nucleic acid may be used to treat (NOGO. The central nervous system (NOGO. The value acident (CVA.)
                       disease, muscure, states which respond to the modura states which respond to the modura
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28-FEB-2000;
06-MAR-2000;
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(BLAT/)
(MCSW/)
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                                                                                                           central nervous system (CNS) injury and cerebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), parkinson's disease, ataxia, Huntington's disease, Creutzfeldt-Jakob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid molecules, e.g., enzymatic nucleic acids and constructs, which down regulate expression of a CD20 gene growth inhibitor gene useful for treating, e.g., lymphoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 88; Page 79; 200pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and central nervous system injury
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Synthetic.
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Parkinson's disease; ataxia; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia;
B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia;
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BLATT L.
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CHOWRIRA B N
                                                                muscular dystrophy, and/or other neurodegenerative hich respond to the modulation of NOGO expression. T
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; 2000US-185516P.
; 2000US-187128P.
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e or neurite
a, leukemia,
                                                                                          disease
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Sequence

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Length 17;

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ABK03622
ID ABK03622
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Best Local Similarity
Matches 15; Conserv
The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGCO).

The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN cleaving NA with a NYN with a NYN triplet), a zinzyme (cleaving RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg^2+. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition
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28-FEB-2000;
06-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury
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(BLAT/)
(MCSW/)
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BLATT L.
MCSWIGGEN J.
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; 2000US-185516P.
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Pred. No. 1.8e
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11-FEB-2000; 28-FEB-2000; 206-MAR-2000; 200

2000US-181797P. 2000US-185516P. 2001WO-US04273

2000US-187128P PHARM

09-FEB-2001;

16-AUG-2001. WO200159103-A2 Synthetic

(RIBO-) (BLAT/) (MCSW/)

MCSWIGGEN J. RIBOZYME

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RESULT 166
ABK03757
ID ABK03757
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AC ABK037
XX
DT 12-WAI
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                           Creutzfeldt-Jakob disease; muscular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                associated with the level of CD20. The treatment may further comprise the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity hes 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1701 TCCAAGAGATAAGCTGA 1717
                                                                                                                                                                                                                                                                                                                                                                 sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 UCCAAGAGACAUGCUGA 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.0%;
76.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 C; 4 G; 3 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 13.8; D
Pred. No. 1.8e
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                           dystrophy; neurodegenerative disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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RESULT 167
ABV788
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ABV788
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ABV788
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DI 03-JAN
XX
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XX

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC CD20 activity of the cell and treat a patient having a condition
CC associated with the level of CD20. The treatment may further comprise the
CC use of one or more therapies. In particular, the CD20 targetting
CC nucleic acid may be used to treat lymphoma, leukaemia, B-cell
CC lymphoma, low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky
CC low-grade or follicular NHL, lymphocytic leukaemia, B-cell
CC immunodeficiency virus) associated NHL, mantle-cell lymphoma (MCL),
CC immunocytoma (IMC), small B-cell lymphocytic lymphoma, immune
CC include acid is used to cleave RNA of the NOGO gene in the presence of a
CC divalent cation that is preferably Mg^2+. Furthermore, the nucleic acid
CC may be contacted with a cell to reduce NOGO activity of the cell and
CC treat a patient having a condition associated with the level of NOGO. The
CC in particular, the NOGO-targetting nucleic acid may be used to treat
CC central nervous system (CNS) injury and cerebrovascular accident (CVA,
Stroke), Alzheimer's disease, demantia, multiple sclerosis (MS),
CC parkinson's disease, ataxia, Huntington's disease, Creutzfeldt-Jakob
CC disease, muscular dystrophy, and/or other neurodegenerative disease
CC states which respond to the modulation of NOGO expression. The
CC present sequence is an amberzyme molecule of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNAzyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN motif) pr an amberzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably MG 20-th Furthermore, it may be contacted with a cell to reduce the contacted with a cell to contacted with a cell to contact the contact the contacted with a cell to contact the co
                                                                                                                            Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placen prostate; skeletal muscle; colon; male infertility; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                 Human HTPL scanning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABV78899 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ŗ
                                                                                                                                                                                                                                                                                                                                                     oligonucleotide SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
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Pred. No. 1.8e
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                          ID 145
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.8e+02;
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down
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RESULT 168
ABV80430/C
ID ABV804
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AC ABV804
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AC ABV804
XX
DT 03-JAN
XX
DE Human
XX
KW Human;
KW Human;
KW male g
KW prosta
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC protein (HTPL, see ABV7875) to ABV78762 and ABB9819 to ABB98520). HTPL

CC has two isoforms, with a few single base pair differences between the

Ct wo. One of the single base pair changes introduces a premature stop

CC shares an overall structure organisation with the Patched protein. The

CC shares an overall structure organisation with the Patched protein. The

CC shares an overall structure organisation with the Patched protein. The

CC important in regulating male germ cell development, and the HTPL is

CC import to human chromosome 10pl2.1. HTPL and its coding sequence are

CC compand to human chromosome 10pl2.1. HTPL and its coding sequence are

CC thorapy and manufacture of a medicament for treatment or prevention of

CC thorapy and manufacture of a medicament for treatment or activity of human

CC foetal liver, bone marrow, brain, kidney, lung, placenta, prostate,

CC skeletal muscle or colon function. HTPL proteins and nucleic acids are

CC clinically useful diagnostic markers and potenial therapeutic agents for

CC male infertility and cancer. The present oligonucleotide was used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                               Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   clinically useful diagnostic markers and male infertility and cancer. The present example from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel isolated human testis expressed Patched like protein useful for identifying agonist and antagonist and specific partners, and for treating subjects having defects in HTPL
                                                                                                                                                                                  Human HTPL scanning
                                                                                                                                                                                                                                                  03-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                        ABV80430
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to human testis expressed Patched like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Page 82; 718pp; English.
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30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                           standard; DNA;
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2001WO-US00667.
2001WO-US00668.
2001WO-US00669.
2001US-0864761.
2001US-0327898.
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                                                                                                                                                                                                                                                  (first entry)
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2001WO-US00664.
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                                                                                                                                                                                     oligonucleotide
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Pred. No. 1.8e+02;
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                                                                                                                                                                                     SEQ ID 1676.
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RESULT 169
ABV09/C
ID ABV09/C
ID ABV09/B
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AC ABV894
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DT 23-DEC
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DE Human
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Human
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                shares an overall structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was mapped to human chromosome 10p12.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
               Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;
                                                                     Human POSHL1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-8 (S for short) compared to HTPL-L (L for long). HTPL become in the compared to HTPL-L (L for long).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel isolated human testis expressed Patched like protein useful for identifying agonist and antagonist and specific partners, and for treating subjects having defects in HTPL
                                                                                                           23-DEC-2002
                                                                                                                                              ABV89409
                                                                                                                                                                              ABV89409 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 5 A; 2 C; 1 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     male infertility and cancer. The example from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Page 283; 718pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-676582/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zhan J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EP1229046-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001;
23-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-JAN-2002; 2002EP-0001167
                                                                                                                                                                                                                                                                                                        2186 ATGTGATGAAAATAGCA 2202
                                                                                                                                                                                                                                                                          17
                                                                                                                                                                                                                                                                                                                                                               Similarity
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2001WO-US00664.
2001WO-US00667.
2001WO-US00667.
2001WO-US00668.
2001WO-US00669.
2001US-0327898.
                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                     scanning oligonucleotide SEQ ID NO 122.
                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                             1.0%;
                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                              <u>.</u>.
                                                                                                                                                                                                                                                                                                                                                             Score 13.8; DB 1;
Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                            Mismatches
             gene
                                                                                                                                                                                                                                                                                                                                                                              Length 17;
             cancer; vaccine;
                                                                                                                                                                                                                                                                                                                                            Indels
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RESULT 170
ABV89410/c
ID ABV894
XX
AC ABV894
XX
AC ABV894
XX
CT 23-DEC
XX
DT 23-DEC
XX
DE Human
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                                                                                                                                                                                                                                                                                                           The invention relates to an isolated SH3 domain (POSH)-like signalling CC protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino cc acids (SI, ABBB399), a sequence having 65% sequence identity to (S1), CC (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. CHuman POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (1) and nucleic acids (II) cencoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL including diagnosing and creating cancer, they useful in the development of vaccines and (II) is useful for measuring and for surveying gene expression and creating care useful for measuring and for surveying gene expression and creating creating cance is that of a scanning oligonucleotide useful in examples of the invention.
                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                        Query Match
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30-JAN-2001;
30-JAN-2001;
                                 23-DEC-2002
                                                               ABV89410
                                                                                             ABV89410 standard;
                                                                                                                                                                                                                                                                                                                                     Note: The present sequence did not form part of the specification, but is based on sequence information by the European Patent Office.
                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 6 A; 4 C; 5 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; SEQ ID NO 122; 60pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression or activity of human POSHL1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-684061/74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shannon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-MAY-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (AEOM-) AEOMICA INC
                                                                                                                                                                                                1564 TCGGCTGAGTCCAGCTC 1580
                                                                                                                                                                                                                                          1 Similarity
15; Conserv
                                                                                                                                                                            TCTGCTGAGTTCAGCTC
                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2001WO-US00664.
2001WO-US00665.
2001WO-US00666.
                               (first entry)
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2001US-0864761.
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2001WO-US00669.
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                                                                                             DNA; 17
                                                                                                                                                                                                                                                       88.2%;
                                                                                                                                                                                                                                                                         1.0%;
                                                                                                                                                                                                                                          <u>.</u>
                                                                                                                                                                                                                                                         Score 13.8; DB 1;
Pred. No. 1.8e+02;
                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                      Length 17;
                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                      printed
supplied to Derwent
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Human POSHL1 scanning oligonucleotide SEQ ID NO

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                                             ABN06320
                                                           RESULT 171
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Best Local (
                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                     protein 1 (POSHI 1) polypeptide (I), comprising a sequence of 730 amino acids (S1, ABB8399), a sequence having 65% sequence identity to (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHI 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHII including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene
                                                                                                                                                                                                                                                            Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied by the European Patent Office.
                            ABN06320 standard; DNA; 17
                                                                                                                                                                                                                                 Sequence 17
                                                                                                                                                                                                                                                                                                                          present sequence is that of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an isolated SH3 domain (POSH)-like signalling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-684061/74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JAN-2001;
23-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-JAN-2002; 2002EP-0001165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (AEOM-) AEOMICA INC
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                                                                                                                                                                                                                                                                                                           the invention.
                                                                                                                                       1563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy;
                                                                                                        17
                                                                                                                                                                     15;
                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; SEQ ID
                                                                                                                            TTCGGCTGAGTCCAGCT 1579
                                                                                                        TTCTGCTGAGTTCAGCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001WO-US00668.
2001WO-US00669.
2001WO-US00670.
2001US-0864761.
2001US-0328205.
                                                                                                                                                                                                                               BP; 7
                                                                                                                                                                     Conservative
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2001WO-US00667.
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                                                                                                                                                                                                                               A; 4 C; 4 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NO 123; 60pp + Sequence Listing; English
                                                                                                                                                                                    88.2%;
                                                                                                                                                                                                  1.0%;
                            ВÞ
                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                          a scanning oligonucleotide useful
                                                                                                                                                                                    Score 13.8; DB 1;
Pred. No. 1.8e+02;
                                                                                                                                                                  Mismatches
                                                                                                                                                                                              Length
                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                            supplied to Derwent
                                                                                                                                                                                                    17
                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                           in examples
                                                                                                                                                                  Gaps
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ABN06320

Query Match

Sequence

17

BP; 3 A;

9 C; 4 G; 1 T; 1.0%;

0 other;

Score 13.8;

DB 1;

Length 17

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The present invention describes a human genome-derived myosin-like CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of CC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 nucleic acids can be used as probes to detect, characterise CC hGDMLP-1 nucleic acids can be used as probes to detect, characterise CC and quantify hGDMLP-1 nucleic acids in samples, as amplification CC substrates, to provide initial substrates for the recombinant engineering CC of hGDMLP-1 protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may CC be used as immunogens to raise antibodies that specifically recognise CC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific CC deficiency in hGDMLP-1 production, and in vaccines or for replacement CC diagnosing a disorder associated with the expression of hGDMLP-1, in CC chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
invention.

N.B. The sequence data for this patent did not form specification, but was obtained in electronic format at ftp.wipo.int/pub/published_pct_sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gu Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human GDMLP-1 17-mer scanning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-179446/23.
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30-JAN-2001;
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27-SEP-2000;
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2001WO-US00664.
2001WO-US00665.
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2001WO-US00667.
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2000US-236359P.
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t directly fr
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                                            from WIPC
                                                                 printed
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88.2%;

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RESULT 172
ABN07681/c
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21-SEP-2000;
27-SEP-2000;
                  of hdDMLP-1 protein variants having desired phenotypic improvements, an for expressing the proteins. The hdDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hdDMLP-1 proteins, as standards in assays used to determine the hdDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hdDMLP proteins, as specifically of hdDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as therapeutic supplement in patients having specific
                                                                                                                                                                            The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gen therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering the property of the provide initial substrates for the recombinant engineering the provide initial substrates for the provide ini
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                                                                                                                                                              substrates, of hGDMLP-1
                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID 7673; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                             myosin-like
                                                                                                                                                                                                                                                                                                                                                                                                              New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human
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replacement
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Matches 15; Conserv
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30-JAN-2001;
30-JAN-2001;
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                                                     New polypeptide, for raising antibodies that proteins, or as specific biomolecule capture surface enhanced laser desorption ionization,
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                     Disclosure;
                                            myosin-like
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04-OCT-2000;
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Pred. No. 1.
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                                                                  recognize hGDMLP-1 probes for
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                                                       comprises
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heart disease;
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The present invention describes

a human genome-derived myosin-like

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ABK26439
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Best Local :
                                                      01-JUN-2000;
30-OCT-2000;
27-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     porphyric herbicide resistance; triazine resistance; disease resistance; modified oil production; modified starch production; waxy starch; altered floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced palmitate production; albino plant; increased stearate production; reduced linolenic acid production;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chromosomal genomic alteration; genome altering oligonucleotide; Po-methyl modification; LNA modification; phosphorothicate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin-B;
                                                                                                                                                                                   01-JUN-2001; 2001WO-US17672.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  photosynthetic
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; 2001US-0818875.
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Pred. No. 1.8e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 part of the printed
t directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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cc phosphorothicate linkages on a terminus, or a combination of any two or combre of these modifications. The oligonucleotides are useful for cdirecting repair or alteration of plant genetic information. The coligonucleotides are particularly useful for creating plants with desired content open and the content of plants or altering amino acid content of plants or conferring amino acid content of plants or conferring amino acid over production), herbicide resistance (e.g. cglyphosate resistance, imidazolinone and sulphonylurea herbicide resistance, porphyric herbicide resistance or triazine resistance), cc disease resistance, modified oil production, modified starch production ce.g. increased starch or production of waxy starch), altered floral comphology (e.g. male-sterile plants) or modified fatty acid content (e.g. reduced palmitate, increased stearate or reduced linolenic acid). Cf he oligonucleotides are also useful for producing albino mutants for the canalysis of photosynthetic processes. This sequence represents a genome creating oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an oligonucleotide for targeted alteration genetic sequence, which comprises a single-stranded oligonucleotide having a DNA domain. The DNA domain has at least one mismatch with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production -
  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                       consist of o-methyl modification, an LNA modification, two or more
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 7; Page 151; 220pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-106307/14
     17
BP; 8
  Α,
     4 C;
     N
     G; 3 T;
     0
       other;
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                S
                                   Matches
                                            Query Match
Best Local
                 1485 CAAGAAGCCAGACTTCA 1501
                                            Similarity
CAAGAAGCTAAACTTCA
                                    Conservative
                                            88.2%;
                                  0;
                                            Score 13.8; DB Pred. No. 1.8e+(
                                    Mismatches
                                            .8e+02;
                                                     Length 17;
                                    Indels
                                   0;
                                   Gaps
                                    0
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RESULT 175
ABK26440/c
Waxy starch
                                                                                      09-APR-2002
                                                                                                     ABK26440 standard; DNA; 17
                                                                             production genome altering oligonucleotide #96.
                                                                                      (first entry)
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Kmiec

EB,

Gamper HB,

Rice

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Kim

06-DEC-2001

WO200192512-A2

Chromosomal genomic alteration; genome altering oligonucleotide; PCR; 88; o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin-B; abiotic stress tolerance; improved nutritional value; hygromycin; primer; amino acid over production; herbicide resistance; glyphosate resistance; imidazolinone herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; triazine resistance; disease resistance; modified oil production; modified starch production; waxy starch; altered floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced paimitate production; albino plant; and plant; albino plant; albino mutant; modified fatty acid content; reduced paimitate production; Solanum tuberosum photosynthetic increased stearate production; acid production

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RESULT 176
ABT35370/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an oligonucleotide for targeted alteration of a CC genetic sequence, which comprises a single-stranded oligonucleotide CC having a DNA domain. The DNA domain has at least one mismatch with CC respect to the genetic sequence to be altered and further comprises CC chemical modifications of the oligonucleotide. The chemical modifications of the oligonucleotide. The chemical modifications CC consist of o-methyl modification, an LNA modification, two or more CC disponsible to these modifications. The oligonucleotides are useful for CC directing repair or alteration of plant genetic information. The CC disponucleotides are particularly useful for creating plants with desired CC phenotypes, e.g. environmental or abiotic stress tolerance, improved CC untritional value (e.g. altering amino acid content of plants or CC conferring amino acid over production), herbicide resistance (e.g. glyphosate resistance, imidazolinone and sulphonylurea herbicide CC disease resistance, modified oil production, modified starch production CC (e.g. increased starch or production of waxy starch), altered floral CC (e.g. reduced palmitate, increased starate or reduced linolenic acid). The oligonucleotides are also useful for producing albino mutants for the CC analysis of photosynthetic processes. This sequence represents a genome CC altering oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUN-2000;
30-OCT-2000;
27-MAR-2001;
                                                                                                          Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease;
             WO2003025175-A2
                                                                              schizophrenia;
human fukutin;
                                                                                                                                                          Tumour suppression
                                                                                                                                                                                               12-JUN-2003
                                                                                                                                                                                                                                                               ABT35370 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 7; Page 151; 220pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified or disease resistance, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kmiec
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                                                                                                                                                                                                                                                                                                                                                                                1485 CAAGAAGCCAGACTTCA 1501
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; 2000US-208538P.
; 2000US-244989P.
; 2001US-0818875.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            production
                                                                                protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; 2 C; 4 G; 8 T; 0 other;
                                                                                                                                                              related
                                                                                                                                                                                                                                                                                                                                                                                                                                 1.0%;
                                                                                             chip; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rice
                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                              human
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                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 13.8; DB 1;
Pred. No. 1.8e+02;
                                                                                                                                                          fukutin oligo SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kim J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 177 ABT39340/c ID ABT393

ABT39340 standard; DNA;

17

ABT39340

12-JUN-2003

(first entry)

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1848 GAAAGACCTTTCTGATC 1864

0;

17 GAAAGACTTTTGTGATC 1

27-MAR-2003

Homo sapiens

fukutin;

gs.

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression;

Tumour suppression related human fukutin oligo

SEQ ID

ö 4977

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sequence that hybridizes to them under highly stringent conditions, or che complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, c.g. as one component of a gene chip, in vitro as (anti) sense reagents, can for production of recombinant polypeptides. Any of the nucleic acids, collypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in gatient samples is useful for diagnosis and/or prognosis of these chips. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression
  Matches
                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a
                                                                                 Sequence 17 BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 150;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polypeptides, antibodies and transfected cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-SEP-2002; 2002WO-IB04208.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel isolated 17 mer nucleic acid sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MOLE-) MOLECULAR ENGINES LAB
l Similarity
15; Conserv
  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amson R,
                   1.0%;
                                                                                 4 C; 2 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       720pp; French
0;
                     Score 13.8;
Pred. No. 1.
    Mismatches
                   .8e+02;
                                      DB 1;
                                      Length 17;
  Indels
0,
Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; prain cancer; coesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; deatrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; deatrexate; gemeitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention.
                                                                                                                                                                                                                                                                          Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;
                                                                                                                                                                                                                                                                                                                         NFKB sub-unit modulating amberzyme substrate #207.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACA09044 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         given in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 615; 720pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polypeptides, antibodies and transfected cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Telerman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-SEP-2002; 2002WO-IB04208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1898 GGAAACACAAGAATATC 1914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention relates to a novel isolated 17 mer nucleic acid sequence, in the specification, a sequence containing at least 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2003-313353/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tuijnder M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 В₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8; DB 1;
1.8e+02;
ches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
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RESULT 179
ABZ64846/c
ID ABZ648
XX
AC ABZ648
XX
DT 21-MAR
XX

ABZ64846;

ABZ64846

standard; RNA; 17

0

21-MAR-2003

(first entry)

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                                                                                                                                                                                                                                                                                                                               The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor (kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme (configuration. The enzymatic nucleic acid molecule is adapted to treat configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A. gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oseophageal, stomach, bladder, pancreatic, correctal, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, methotrexate, colorentes or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, chesity, astchman, crohn's disease, diabetes,
                                                                                    Matches
                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-AUG-1994;
07-DEC-1992;
18-MAY-1994;
23-DEC-1996;
                                                                                                                                                                                                                                               rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or sepsis, allergic airway inflammation, inflammatory bowel disease or
                                                                                                                                                                      Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2002177568-A1.
                                                                                                                                                                                                              infection. This sequence represents the substrate enzymatic nucleic acid molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Page 55; 72pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-340953/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stinchcomb DT, Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (STIN/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-NOV-2002.
                                                                             Local Similarity
hes 15; Conserv
                                          2004 AGCCCGGAGGCCACCCG
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DRAPER K G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STINCHCOMB D
  AGCCCAGAGGCCCCCCG
                                                                                                                                                                    BP; 3 A;
                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92US-0987132.
94US-0245466.
96US-0777916.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94US-0291932
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                                                                                                      1.0%;
                                                                                                                                                                    9 C; 5 G; 0 U; 0 other;
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                                          2020
17
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                                                                                                      Score 13.8;
Pred. No. 1.
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                                                                                    Mismatches
                                                                                                      1.8e+02;
                                                                                                                              DB 1;
                                                                                                                          Length 17;
                                                                                    Indels
                                                                                  <u>,,</u>
                                                                                    Gaps
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Human HER2 DNAzyme substrate #303

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RESULT 180
ABZ65209
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-R human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66530 - ABZ66531 represent substrate/target sequences for the human ribozymes of the invention.
                                                                                         Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-MAY-2001; 2001US-294140P.
06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
05-DEC-2002.
                               WO200297114-A2
                                                           Homo sapiens
                                                                                                                                                         Human HER2 DNAzyme
                                                                                                                                                                                        21-MAR-2003
                                                                                                                                                                                                                                                     ABZ65209 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 5 A; 3 C; 6 G; 3 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel short interfering RNA (siRNA) nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 138; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-140484/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAY-2002; 2002WO-US16840.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-DEC-2002.
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                                                                                                                                                                                                                                                                                                                                                               1920 TCTTCTTGGAGCCTGCA 1936
                                                                                                                                                                                                                                                                                                                                   17
                                                                                                                                                                                                                                                                                                                                                                                                15;
                                                                                                                                                                                                                                                                                                                                 TCTTCTTGCAGCCAGCA 1
                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                       (first entry)
                                                                                                                                                         substrate #666.
                                                                                                                                                                                                                                                                                                                                                                                                             1.0%;
                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                               Score 13.8; DB 1; Length 17; Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                              0
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RESULT 181
AAQ49011
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-MAY-2001;
06-JUN-2001;
10-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SBP; specific binding pair members; antibody; RGDP; replicable genetic display package; recombination; PCR. polymerase chain reaction; ss.
Griffiths AD,
                                                                                                                                              24-MAR-1992;
15-MAY-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Multimeric (SBP) antibody chain primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
                                                                                                                                                                                                                                   24-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                       WO9319172-A1
                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ49011 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 4 A; 5 C; 5 G; 3 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 145; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-140484/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcswiggen
                                                         (CAMB-) CAMBRIDGE ANTIBODY TECHNOLOGY. (MEDI-) MEDICAL RES COUNCIL.
                                                                                                                                                                                                                                                                                             30-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2269 CCAGTCAAGTGGATGGC 2285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CCCAUCAAGUGGAUGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; 2001US-294140P.
; 2001US-296249P.
; 2001US-318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      relates to a novel short interfering RNA (siRNA) nucleic
                                                                                                                                              92GB-0006318.
92WO-GB00883.
                                                                                                                                                                                                                                      93WO-GB00605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>س</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 13.8; DB 1;
Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB · 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               H-Ras, N-Ras,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
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Johnson KS,

Smith AJH,

Waterhouse

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Query Match

Length

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                                                    The sequences given in AATO5636-39 are primers which were used construction of a mutated factor VIII molecule. The amplified encodes a molecule in which Arg 562 is substituted for Ile. The mutation occurs in the cleavage site for activated protein C (A confers resistace to APC cleavage. The novel factor VIII based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The primers (AAQ48987-Q49045) are used in the amplification of Kappa and lambda-chain genes of various antibodies. These genes are then recombined into the same replicon, resulting in very diverse libraries of antibody chains, e.g. from unimmunised donors. It is also useful for chain shuffling, mutagenesis, humanising and CDR imprinting. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                             substitution; factor V; acti resistance; thrombo-embolic
                                                                                                                                                                                                                                                                                                                                                                                                                                               Primer; amplify; polymerase chain reaction; PCR; diagnosis; intron 10; substitution; factor V; activated protein C; APC; cleavage site; resistance; thrombo-embolic disease; coagulation cascade; ss.
                                      confers resistace to APC cleavage. The nov
                                                                                                                                                                                     Activated
                                                                                                                                                                                                                  WPI; 1995-383004/49
                                                                                                                                                                                                                                                                                                     22-APR-1994;
                                                                                                                                                                                                                                                                                                                                21-APR-1995;
                                                                                                                                                                                                                                                                                                                                                              02-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                         WO9529259-A1
                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Primer F8-547S sense to bases 547-565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT05636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT05636 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 7 A; 6 C; 3 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 58; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1993-320739/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Winter GP
                                                                                                                                                                                                                                                                          (BLOE-) STICHTING CENT LAB VAN DE BLOEDTRANSFUSI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     odn. of specific binding pair members, e.g. antibody chains display on surface of replicable genetic display packages
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1673 AACTTCCAGAGGACCCA 1689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                       protein C resistant mutant factors V or VIII - useful for and treating disorders in the blood coagulation cascade
                                                                                                                                                                     and treating
                                                                                                                                          Page 23; 48pp; English.
                                                                                                                                                                                                                                             Van Mourik JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                     94EP-0201116
                                                                                                                                                                                                                                                                                                                                  95WO-NL00149
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                                                                                                                                                                                                                                             Voorberg JJ
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     No. 1.
                                        for activated protein C (APC) which The novel factor VIII based protein orders in the blood coagulation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of factor VIII cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; DB 1;
L.9e+02;
hes 2;
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Sequence 18

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Sequence 18

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RESULT 183
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Matches
                                       some or all of the following elements: purification tag:proteinase site:ECM binding site:proteinase site:TGF-beta. TGF-beta promotes wound healing, and the fusion protein can be used to reduce surgery recovery time and in the preparation of artificial skin. The inclusion of a purification tag facilitates purification of the fusion protein. The proteinase site is included to permit cleavage and release of the purification tag after purification if desired. The extracellular matrix binding site facilitates delivery of the fusion protein to the desired site of action. Delivery of the TGF-beta to the site to be treated reduces the amount of TGF-beta required to be administered to be effective and reduces the concentration of circulating TGF-beta which may result in undesirable effects.
                                                                                                                                                                                                                                                                                                                                              A novel transforming growth factor-beta (TGF-beta) fusion protein comprises a purification tag and a TGF active fragment. The present sequence encodes a specifically claimed purification tag.

Additionally, the fusion protein may comprise proteinase-sensitive highest sites and binding domain so the protein sequence may contain the protein the protein sequence may contain the protein the protei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT71737
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT71737;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 40; 59pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Prepn. of transforming growth factor-beta fusion protein - useful reduce surgery recovery time and to prepare artificial skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cheung DT, Hall
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NIMN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Transforming growth factor-beta fusion protein; wound healing; artificial skin; surgery recovery time; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-AUG-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CHEU/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MULT/)
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TUAN T.
WU L.
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HALL F
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Pred. No. 1.9e+02;
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RESULT 184
AAV48746/c
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Best Local
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                                                                                                                                                                                                                                                           ErbB-2 gene. Of these, only oligonucleotides AAV48709-91 resulted in significant redcution in ErbB-2 protein expression, while oligonucleotides AAV4872-886 had little effect. The oligonucleotides exemplify the invention. The specification describes oligonucleotides that contain 8-30 nucleotides, which contain at most 8 nucleotides that can each form three hydrogen bonds to cytosine; do not contain four consecutive nucleotides able to form three H-bonds each to four consecutive cytosine; do not contain two sequences of three consecutive nucleotides each able to form three H-bonds to three consecutive consecutive and the ratio between residues able to form two H-bonds each (2R) or three such bonds (3R) is given by ZR/JR = 0.33-0.72. The oligonucleotides are used to modulate expression of genes, particularly the genes for p53, ErB-2, junB, junD, TGF-beta 1 or beta 2 to control proliferation of primary cell cultures (e.g. bone marrow stem, liver or by the contain the contains of primary cell cultures (e.g. bone marrow stem, liver or by the contains of primary cell cultures (e.g. bone marrow stem, liver or by the contains of primary cell cultures (e.g. bone marrow stem, liver or by the contains of primary cell cultures (e.g. bone marrow stem, liver or by the contains of primary cell cultures (e.g. bone marrow stem, liver or by the contains of primary cell cultures (e.g. bone marrow stem, liver or by the contains of the 
                                                                                                                                                                                                 kidney cells, osteoclasts, osteoblasts and/or keratinocytes). The oligonucleotides can also be used to analyse function of proteins (by altering their expression or activity) and therapeutically, e.g. in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Preparation of antisense oligo:nucleotide(s) which lack long runs consecutive guanosine or inosine - and have specific ratio of residues able to form two or three hydrogen bonds, have greater activity and reduced toxicity, used therapeutically or to modulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                       Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV48709-886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 10; Fig 6a; 286pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Brysch W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-JAN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ErbB-2; antisense oligonucleotide; modulate; gene expression; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ErbB-2 gene antisense oligonucleotide ErbB-2-38.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BIOG-)
                                                                   Local
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  2317 CATCAGAGTGATGTCTG 2333
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                                            al Similarity
15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-886 represent antisense oligonucleotides directed against the
gene. Of these, only oligonucleotides AAV48709-91 resulted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schlingensiepen K;
                                                                                                                                BP; 4 A; 6 C; 4 G; 4 T; 0 other;
                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                              1.0%;
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                                                                 Score 13.8; DB Pred. No. 1.9e+(
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Pred. No. 1.
                                                                                                                                                                              TGF)
                                            Mismatches
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                                                                                                                                                                          for stimulating the immune
                                                              8; DB 1;
1.9e+02;
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                                            <u>ب</u>
                                                                                   Length 18;
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                                            Indels
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                                                                                                                                                                              system
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RESULT 186
AAV99371/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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Best Local
                                                                                                                  cDNA encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to new influenza antigens that comprise a fusion product of at least the extracellular part of a conserved influenza membrane protein (or a functional fragment) and a presenting carrier. The membrane protein consists of the extracellular part of the influenza matigens can be used in the preparation of a vaccine against influenza for humans and animals, e.g. pigs and horses. The vaccines can be direct vaccines, e.g. vaccines containing the fusion
                                                                   Proteinase site; bone morphogenetic fusion protein; bone binding site;
                                                                                                                                                                                                                        AAV99371;
                                                                                                                                                                                                                                                                          AAV99371 standard; cDNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Fig 29; 100pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>X</u>
                       active
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fiers W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RT-PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Influenza antigen; fusion product; extracellular; membrane protein;
M2 protein; vaccine; human; animal; pig; horse; RT-PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   1821 GAAGATGTTGAAAGATG 1837
                                            morphogenetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17
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15; Conservat
                       fragment;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  or indirect,
                                                                                                                                                                                                                                                                                                                                                                                                    GTAGATATTGAAAGATG 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                      (first entry)
                                                                                                                     a peptide comprising a purification tag.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                    wound healing;
                                            protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 C; 6 G; 5 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Neirynck S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ٥,
                                         transforming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 13.8; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 2;
                       bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              얁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 other;
                    ming growth growth; pur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ø
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.9e+02;
2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              conserved
                       purification tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           comprising a
/ed influenza
                                                 factor beta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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membrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence encodes a peptide comprising a purification tag that was used in the creation of the bone morphogenetic fusion proteins of the invention. The bone morphogenetic fusion protein may contain some or all of the following elements: a purification tag, a proteinase site, an ECM/bone binding site, a second proteinase site, and a bone morphogenetic protein active fragment. The fusion proteins of the invention also includes proteins that have transforming growth factor beta active fragments instead of bone morphogenetic protein active
                    protein trafficking; cell surviva. DNA synthesis; apoptosis; neurite glucose transport; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-059875/05.
                                                       Phosphatidylinositol 3-kinase; PI3K; p85; p110; heterodimer; hormone; growth factor; receptor; antisense; inhibition; expression; diagnosis; modulation; growth factor mediated cell transformation; mitogenesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HANB/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDS
                                                                                                                      Antisense
                                                                                                                                             19-DEC-2000
                                                                                                                                                                                            AAA52031 standard; cDNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                     Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                        rragments. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 37; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      wound healing or bone growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New bone morphogenetic fusion tag and a bone morphogenetic a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hall FL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-DEC-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (WULL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SHOR/)
                                                                                                                                                                                                                                                                    1879 GAGATGATGAAGATGAT 1895
                                                                                                                                                                                                                                                       18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ) NIMNI M E.
) SHORS E C.
) WU L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HALL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW84203.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HAN B.
                                                                                                                                                                                                                                                       GTGATGATGATGAT
                                                                                                                    oligonucleotide directed against PI3K p85 subunit.
                                                                                                                                                                                                                                                                                                                                                                            . The bone morphogenetic fusion proteins can be used for wound healing or bone growth.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Han B,
                                                                                                                                                                                                                                                                                                                                                     BP; 6
                                                                                                                                                                                                                                                                                                      Conservative
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ር.
                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nimni ME,
                                                                                                                                                                                                                                                                                                                                                     A; 7 C; 0 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                  1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "encodes a purification tag"
                              survival; cell proliferation; neurite outgrowth; insulin-stimulated
                                                                                                                                                                                                                                                         N
                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                     Score 13.8; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shors EC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   active fragment, used for enhancing
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                                                                                                                                                                                                                                                                                                     2;
                                                                                                                                                                                                                                                                                                                             Length 18;
                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                     0;
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RESULT 188 AAA48791/c ID AAA487 XX

AAA48791 standard;

DNA;

18

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17

TTGGAAGAAGACTTGAA

Matches Best Local

15; Conserv

Conservative

<u>,</u>

Mismatches

Indels

0

Gaps

0

2530 TTGGTAGAAGACTTGGA 2546

Human; G-alpha-16; G protein; cytostatic; hyperproliferative disorder; cancer; inflammation; infection; antisense inhibition; ss.

cancer; inflammation; infection; antisense

Human G-alpha-16 antisense oligonucleotide ISIS# 20848

08-SEP-2000

(first entry)

WO200032817-A1

Homo sapiens.

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cc and oncogene products. PIJKS act as downstream effectors of these creeptors, are recruited upon receptor stimulation and mediate the cc activation of second messenger signaling pathways. The PIJ kinase cray consists of a 110kD catalytic subunit (p10) associated with cc an 85kD regulatory subunit (p85) and it is through the SH2 domains co f the p85 subunit that the enzyme associates with the membrane cc bound receptors. PIJKs have been implicated in many cellular cc activities including growth factor mediated cell transformation, cc mitogenesis, protein trafficking, cell survival and proliferation. CC mitogenesis, protein trafficking, cell survival and proliferation. CC places transport. Antisense compounds directed against PIJK p85 and cc which inhibit its expression are useful as diagnostics and research cc reagents, and as a component of kits, which can be used for detecting ct the level of pIJK p85 in a sample. The compounds may be administered to be treated by modulating the expression of PIJK p85. The compounds may be administered can be treated by modulating the expression of PIJK p85. The compounds may cc further be useful prophylactically, e.g., to prevent or delay confection, inflammation or tumour formation. The target site of the pIJK p85 subunit (See GENESEQ record AAA52007).
                     Query Match
                                                               Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense compounds targeted to the coding region of human phosphatidylinositol 3-kinase (pI3K) p85 and inhibiting pI3K p85 expression, useful for treating disorders associated with pI3K p85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 with the cytoplasmic domain of hormone and growth factor receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The phosphatidylinositol 3-kinases (PI3Ks) family of heterodimeric lipid kinases that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-AUG-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cowsert LM;
                                                               BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Column 39; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0344521.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0344521.
1.0%;
                                                               5 C; 1 G; 8 T; 0 other;
    Score 13.8; DB 1;
Pred. No. 1.9e+02;
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                       Length 18;
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RESULT 189
AAA23501
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Best Local S
Matches 15
24-AUG-1998;
24-AUG-1998;
09-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence is an antisense oligonucleotide used to modulate expression of G-alpha-16. G-alpha-16 is a human G protein which interacts differentially with several receptor types including members of the opicid and chemokine receptor families. A series of antisense oligonucleotides have been designed to target different regions of the human G-alpha-16 RNA. They may be used to inhibit the expression of
                                                                                                                                                                                             Human; secreted protein; cancer; tumour; cardiovascular disorder; blood disorder; haemophilia; autoimmune disease; diabetes; inflammation; infection; fungal; bacterial; viral; HIV; allergy; arthritis; neurodegenerative disease; asthma; contraceptive; hybridisation probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G-alpha-16 in human cells and tissues and thus to treat diseases associated with G-alpha-16, such as hyperproliferative disorders, especially cancer. Infections, inflammation or tumour formation car be prevented or delayed. The compounds can be used in research and diagnostics in sandwich and other assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A new antisense compound for inhibiting the expression of human G-alpha-16 and treating, preventing or delaying infections, inflammation or hyperproliferative disorders such as cancer -
                                                                                                                                                                                                                                                                                                            19-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Note: The sequence has a phosphorothicate backbone and may be either an oligodeoxynuclectide or a chimeric oligonuclectide containing 2'-methoxyethyl (2'-MOE) wings and a deoxy gap. The ISIS number given above corresponds to the oligodeoxynuclectide sequence
                                                            24-AUG-1999;
                                                                                            02-MAR-2000
                                                                                                                         WO200011015-A1
                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                        Clone vq1_1 hybridisation probe,
                                                                                                                                                                                                                                                                                                                                          AAA23501;
                                                                                                                                                                                                                                                                                                                                                                     AAA23501 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 3; Page 73; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1354 CCAGCGCCTGGAAGAGA 1370
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                                                                                                                                                                                                                                                                                                          (first entry)
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98US-0097638.
98US-0097659.
98US-0099618.
                                                            99WO-US19351.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 7 C; 4 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 13.8; DB 1;
Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                             SEQ ID NO:119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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partially secreted, i.e., transmembrane proteins. The proteins of the invention may exhibit one or more activities selected from the following cytokine activity; cell proliferation; differentiation; immune modulation; haematopoiesis regulation; tissue growth activity; haematopoiesis regulation; tissue growth activity; activin/inhibin activity; chemotactic/chemokinetic activity; haemostatic and thrombolytic activity; anti-inflammatory activity; and tumour inhibition activity. The proteins may be administered to patients as vaccines, and the nucleotides may be used as near of constitutions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-NOV-1998;
23-DEC-1998;
23-DEC-1998;
23-AUG-1999;
                                                                                                                                                                                        vaccines, and the nucleotides may be used as part of a gene therapy regime. Diseases or conditions that may be treated using the proteins or nucleotides of the invention include autoimmune diseases; genetic disorders; haemophilia; cardiovascular diseases; cancer; bacterial, fungal and viral infections, especially HIV; multiple sclerosis; rheumatoid arthritis; pulmonary inflammation; Guillain-Barre syndrome; insulin dependent diabetes mellitus; and allergic reactions such as asthma and anaemia. They may also be used for treating wounds, burns, ulcers, osteoporosis, osteoparthritis, periodontal diseases, Alzheimer's disease, Parkinson's disease, Huntington's disease and amyotrophic lateral sclerosis (ALS). Proteins with activin/inhbin activity may additional activity may additional disease.
                                          Sequence 18
                                                                                                         additionally be useful as contraceptives. Nucleic acid sequences of the invention may be used in chromosome mapping, and as a source of diagnostic primers and probes. Sequences AAA23463-A23502 represent hybridisation probes which may be used to isolate the cDNA clones of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to 40 human secreted proteins (AAY9498 and cDNA sequences encoding them (AAA23423-A23462). The secreteins of the invention include those that are thought to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 345;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-224657/19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ALPH-) ALPHAGENE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and
                                            BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or transmembrane proteins and polynucleotides encoding for treating neurodegenerative disorders, autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98US-0102092.
98US-0109978.
98US-0113645.
98US-0113646.
99US-0379246.
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                                          5 C; 6 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           357pp; English.
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Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (AAY94981-Y95020),
The secreted
                                                                                                              clones of the
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5 Matches Query Match Best Local : Local Similarity les 15; Conserv 1493 CAGACTTCAGCAGCCAG Conservative 1.0%; 1509 0; Score 13.8; DB 1; Pred. No. 1.9e+02; Mismatches Indels 0, Gaps 0

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CAGACATGAGCAGCCAG

17

RESULT 190 AAZ59167/c 20-APR-2000 AAZ59167 Fusion protein; Bacillus; Hexa(his) oligonucleotide for MWPsp-MWPmp10-(His)6-linker-Met-ProIns. protease; standard; DNA; 18 (first entry) PCR primer; cell wall protein;

promoter; cleavage site;

JP11341991-A

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RESULT 191
AAZ59168
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a DNA construct encoding a fusion protein comprising a Bacillus species cell wall protein fused to a cleavage peptide and a heterologous protein. The fusion construct is placed downstream of a Bacillus species promoter sequence. This sequence represents an oligonucleotide coding for the (His)6 part of the construct WMPsp.WMPmp10-(His)6-Linker-Wet-ProInsulin. This construct comprises the Bacillus brevis middle wall protein mp10 linked to the
                                                                                                                                                                                           14-DEC-1999.
                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A DNA coding a new fused
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sato S,
        Example 1; Page 8; 43pp; Japanese
                              A DNA coding a new fused protein through its expression -
                                                                WPI; 2000-101697/09
                                                                                      Sato S,
                                                                                                                                             31-MAR-1998;
                                                                                                                                                                   30-MAR-1999;
                                                                                                                                                                                                                JP11341991-A.
                                                                                                                                                                                                                                                           TEV protease; PCR
                                                                                                                                                                                                                                                                      Fusion protein; Bacillus; cell wall protein; promoter; cleavage site;
                                                                                                                                                                                                                                                                                           Hexa(his) oligonucleotide for MWPsp-MWPmp10-(His)6-linker-Met-ProIns.
                                                                                                                                                                                                                                                                                                                  20-APR-2000
                                                                                                                                                                                                                                                                                                                                         AAZ59168;
                                                                                                                                                                                                                                                                                                                                                           AAZ59168 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 6 A; 7 C; 0 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proinsulin protein via a cleavable linker sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 8; 43pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          through its expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-MAR-1999;
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                                                                                                              (UDAK/)
                                                                                                                         (ITOH-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UDAK/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                           1879
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                                                                                                            ITOHAM FOODS INCUDAKA S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      UDAKA S.
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                                                                                      Higashikuni N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Higashikuni N,
                                                                                                                                                                                                                                                                                                                                                                                                                      GTGATGATGATGAT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                          GAGATGATGAAGATGAT 1895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                           primer; ss.
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                                                                                       Kudo
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Pred. No. 1.9e+02;
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                                          and preparation of a useful peptide
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AAZ59072
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         NNA complex in presence of the test compound and comparing the result with a standard. The oligonuclectides AAZ59070-Z59071 anneal to form a double stranded oligonuclectide containing the HIV-1 trans-activation regulatory region (TAR) to which the HIV-1 Tat protein binds. The complex is labelled with 6-carboxyfluorescein and is used as a target for the binding of a labelled ADP-1 protein. Detection of the complex is by fluorescence resonance energy transfer (FRET). The method is used to identify compounds that interfere with interaction between the target RNA and ligands or proteins. Compounds that are identified are potentially useful for treating infections (viral, bacterial or fungal), cancer and autoimmune diseases. The compounds are preferably directed to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a DNA construct encoding a fusion protein comprising a Bacillus species cell wall protein fused to a cleavage peptide and a heterologous protein. The fusion construct is placed downstream of a Bacillus species promoter sequence. This sequence represents a reverse oligonuclectide coding for the (His) 6 part of the construct MMPsp.MMPmp10-(His)6-Linker-Met-ProInsulin. This construct comprises the Bacillus brevis middle wall protein mp10 linked to the proinsulin protein via a cleavable linker sequence.
                                                                                                                                                                                                                        The invention relates to a method of determining if a compound binds to a target RNA by treating a test compound with a reporter (R) labelled with a donor or acceptor group and labelled target RNA, labelled with the complementary donor or acceptor group, and measuring the fluorescence from fluorescent groups associated with a compound:target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 5 A; 0 C; 7 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                              Examples; Page 31; 82pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUN-1998;
02-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus type 1..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     autoimmune disease; trans-activation regulatory region; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV-1 TAR oligonucleotide target sequence
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                                                                                                                                                                                                                                                                                                                                                                                 Identifying compounds that bind to target RNA, potentially useful treating infections, tumors and autoimmune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-097545/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RIBO-) RIBOTARGETS LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antiviral; antibacterial; antifungal; anticancer; detection; TAR; RRE; fluorescence resonance energy transfer; tat; HIV-1; Rev response element;
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99GB-0004790.
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                                                                                                                                                                                                                                                                                                                                              English.
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Pred. No. 1.
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1.9e+02;
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RESULT 193
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                                                                                                                                                                   This invention describes novel antisense oligonucleotides (I) capable of inhibiting expression of human EGR-1 (early growth response-1). The products of the invention have anti-inflammatory, cytostatic and antiviral activity. (I) was tested for its effects on EGR-1 mRNA levels by real-time polymerase chain reaction (PCR), results indicated that 60% inhibition was achieved. When (I) was modified by 2'-O-methoxyethyl substitution of the first 4 and last 4 residues, and by replacing any C in these flanking regions with 5-methyl-C, the degree of inhibition was increased to 71%. (I) is used to inhibit expression of EGR-1 in cells and tissues in vitro, for research or diagnosis, e.g. detecting EGR-1 encoding nucleic acid. (I) may also be used to treat or prevent and these thanking particularly vital infections, inflammation and these and the prevent and the seases, particularly vital infections.
                                                                                                                                                                                                                                                                                                                                                                                              Antisense oligonucleotides that inhibit expression of human early growth response-1, useful for diagnosis, treatment and prevention tumors, inflammation and infection -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EGR-1; early growth response 1; antisense; inhibition; human; primer; anti-inflammatory; cytostatic; antiviral; detection; diagnosis; viral infection; inflammation; tumor; ss.
                                                                                                                 Sequence 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human EGR-1 DNA antisense primer #24161.
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                                                                                                                                              human
                         1924 CTTGGAGCCTGCACACA 1940
                                                                                                                                                             tumors.
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                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                 15; Column 37-38; 31pp; English.
 CCTGGAGCCTGCACCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGCCAGAUUUGAGCAGC 17
                                                                                                                                              EGR-1 protein.
                                                                                                                 BP; 2
                                                                                                                                                          AAZ44124-Z44169 represent antisense
                                                        Conservative
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                                                                                                               A; 4 C; 9
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                                                     Score 13.8; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 2;
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Pred. No. 1.9e+02;
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                                                                                                                                                          primers used to inhibit
                                                                                 Length 18
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RESULT_194
AAZ36560
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Best Local
                                                                                                                                                                                                                                                                                                      uses sets of peptide nucleic acid (PNA) probes in hybridisation reactions. The method comprises using at least 2 sets of hybridisation probes, where at least one set comprises one or more PNA probes capable of hybridising to specific nucleic acid sequences related to a potential aberration in a chromosome. The methods can be used for the detection of chromosome aberrations. They can be used for the diagnosis of disorders and diseases related to chromosomal aberrations or abnormalities such as e.g. haemapoietic malignancies, cancers and inborn constituel diseases. The method may be used for detecting viral sequences and their localization in the chromosome. In plant biology, the methods can be used for monitoring the efficiency of transferring herbicide resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ36520-41 (set 1) and AAZ36542-61 (set 2) represent two sets of probes which flank each site of the major breakpoint region (mbr) of the MLL-1 gene. The MLL gene is associated with acute lymphoblastic leukaemia-1 (ALL-1). The probes are selected from the lower strand of the MLL-1 gene, and so hybridise to the upper strand. The probes are used to demonstrate the method of the invention. The specification describes a method for the detection of chromosome aberrations in eukaryotic samples
                                                                                                                                                                                                                              Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detection of chromosome aberrations, used for detecting diseases and disorders, infections, and plant alterations related to e.g. herbicide
                                                                                                                                                                                                                                                                                       genes to a plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 40; 63pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acute lymphoblastic leukaemia-1; ALL-1; probe; peptide nucl haemapoietic malignancy; cancer; inborn constituel disease; herbicide resistance gene; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    resistance
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N
                                          TGTGGGAGATCTTCACT
   TGTGGGAGATGTTGACT
                                                                                                                                                                                                                              BP; 3
                                                                                                                  Conservative
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                                                                                                                                                                                                                              2 C; 7 G; 6 T; 0 other;
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                                                                                                                                          Score 13.8; DB 1;
Pred. No. 1.9e+02;
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                                                                                                                                                                  Length 18;
                                                                                                                  Indels
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RESULT 195 AAF74454

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RESULT 196
AAF74457/c
X E X E X E X E X E X E
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                                                                                                                                                                                                                                                                                                        The present invention describes isolated nucleic acids encoding secreted polypeptides, designated PROX polypeptides (i.e. a PRO polypeptide where X is an integer from 1 to 17). PROX polypeptides have cytostatic, immunomodulatory and reproduction activities, and can be used in gene therapy, and as PROX antagonists and PROX agonists. PROX polypeptides, nucleic acids and antibodies are useful in the manufacture of a medicament for treating a syndrome associated with a PROX-associated disorder, e.g. a cell proliferation and/or differentiation aldorder (e.g. pre-clampsia). They are also used for screening for a modulator of activity or of latency or predisposition to a PROX-associated disorder. AAF74432 to AAF74448 encode the specifically claimed human PROX polypeptides PRO1 to PRO17 given in AAB70531 to AAB70547. The present reguence represents a primer used in an example from the present
                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; PRO; PROX; cytostatic; immunomodulatory; reproduction; gene therapy; cell proliferation; differentiation disorder; cancer; immune associated disorder; gestational disease; pre-clampsia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleic acids encoding secreted polypeptides, designated PROX polypeptides, useful for treating a syndrome associated with PROX-associated disorder, e.g. cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-AUG-1999; 99US-0148433.
10-AUG-2000; 2000US-0148433.
            Human PRO2 gene-specific sequencing primer
                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-MAY-2001
                                            09-MAY-2001
                                                                                                      AAF74457 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-147509/15.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CURA-) CURAGEN CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF74454 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                       invention.
                                                                                                                                                                                                               1481 CGACCAAGAAGCCAGAC 1497
                                                                                                                                                                                                                                             15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Page 119; 166pp; English.
                                                                                                                                                                                  CTACCAAGAAGCCAGCC 18
                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fernandes E;
                                                                                                                                                                                                                                                                                                       A; 8 C; 3 G; 1 T; 0 other;
                                                                                                                                                                                                                                                         1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequencing primer
                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                            Score 13.8; D
Pred. No. 1.9e
0; Mismatches
                                                                                                                                                                                                                                                          8; DB 1;
1.9e+02;
              SEQ ID NO:43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID
                                                                                                                                                                                                                                             2
                                                                                                                                                                                                                                                                          Length 18;
                                                                                                                                                                                                                                             Indels
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RESULT 197
AAF56051
ID AAF560
XX AAF560
XX AAF560
XX IB-APR
DT 18-APR
DX HBV DN
XX HBV; h
KW HBV; h
KW mutati
XX mutati
XX HPati
XX HPati
XX HPati
XX HPATI
PN WC2001
XX HPJAN
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Best Local S
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immune associated disorder; gestational disease; pre-clampsia;
PCR primer; sequencing primer; ss.
                                                                                                                                                    HBV; hepatitis B virus; DNA polymerase mutation detection; probe; ss.
                                                                                                                                                                                                                                HBV DNA polymerase gene M552V/I mutation probe HBPr463
                                                                                                                                                                                                                                                                                                                                   AAF56051;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-AUG-1999; 99US-0148433.
10-AUG-2000; 2000US-0148433.
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gene therapy; cell proliferation; differentiation disorder; c
                                                      WO200104358-A2
                                                                                                     Hepatitis B virus
                                                                                                                                                                                                                                                                                 18-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                    AAF56051 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 119; 166pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1481 CGACCAAGAAGCCAGAC 1497
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTACCAAGAAGCCAGCC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.0%;
llarity 88.2%;
Conservative
                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fernandes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 C;
                                                                                                                                                                                                                                                                                                                                                                                    18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 13.8; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 other;
                                                                                                                                                                           gene; anti-HBV drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                   resistance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>,,</u>
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18-JAN-2001

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AAD40986/c
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Best Local S
Matches 15
                                                                                                                                                                                                                                                          Human; antisense; PI3K p85; obesity; type 2 diabetes; cancer; tumour; prophylaxis; hyperproliferative condition; infection; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is a probe used in a method for monitoring anti-hepatitis B virus (HBV) drug resistance in a patient by genetic detection of any one of mutations L528M, b552V/I and/or V/L/M555I in HBV DNA polymerase in a biological sample from the patient. The method is useful in the field of genetic detection of anti-HBV drug resistance during HBV therapy. The method is rapid, reliable and
                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Monitoring anti-HBV drug resistance by genetic detection of mutations in DNA polymerase of HBV in patient's sample, involves hybridizing the polynucleic acids of the sample with a probe and detecting the hybrid
         modified_base
                                         modified_base
                                                                                                                                   modified_base
                                                                                                                                                                             modified_base
                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                               therapy; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                            Human PI3K p85
                                                                                                                                                                                                                                                                                                                    30-OCT-2002
                                                                                                                                                                                                                                                                                                                                          AAD40986;
                                                                                                                                                                                                                                                                                                                                                                 AAD40986 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 2; Page 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-138370/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stuyver L, Maertens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-JUL-1999;
13-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (INNO-) INNOGENETICS NV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                              2267 TTCCAGTCAAGTGGATG 2283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15;
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                                                                                                                                                                                                                                                                                                                                                                                                                        TTTCAGTCATGTGGATG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 4 A; 2 C; 5 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000WO-EP06306
                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                           antisense oligonucleotide ISIS #28034.
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99US-0143546
          /mod_base=
                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                             /note= "Phosphorothioate backbone"
                                                                                                                                                       mod_base= OTHER
                               /*tag= d
                                                                                                'mod_base= OTHER
note= "2'-methoxyethyl nucleotides"
                                                      notes
                                                                                                                        *tag= .b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              64pp; English.
                                                                                                                                                                                                                                                                                                                    entry)
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                                                                  base=
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                                                      -methoxyethyl nucleotides"
                     m5c
                                                                  OTHER
                                                                                                                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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Pred. No. 1.9e+02;
0; Mismatches 2;
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RESULT 199
ABL30659/c
ID ABL306
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to antisense compounds targetted to a nucleic acid molecule encoding PI3K p85 to inhibits its expression. Antisense compounds of the invention are used for treating obesity. Type 2 diabetes and hyperproliferative condition e.g. cancer. They may also be useful prophylactically, e.g. to prevent or delay infection, inflammation or tumour formation. Antisense compounds either alone or in combination with other antisense compounds or therapeutics can be used as tools in differential and/or combinatorial analyses to elucidate
           Inoko H,
                                     (NISN)
                                                                         01-JUN-2000; 2000JP-0164798
                                                                                                 01-JUN-2001; 2001WO-JP04662.
                                                                                                                          06-DEC-2001.
                                                                                                                                                     WO200192572-A1.
                                                                                                                                                                                                   Human; human leukocyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.
                                                                                                                                                                                                                                         Human HLA genotyping oligonucleotide
                                                                                                                                                                                                                                                                     21-MAR-2002
                                                                                                                                                                                                                                                                                             ABL30659;
                                                                                                                                                                                                                                                                                                                     ABL30659 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 4 A; 5 C; 1 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotide targetted to human PI3K p85 DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         expression patterns of a portion or the entire complement of genes expressed within cells and tissues. They are commonly used as research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense compounds targeted against polynucleotides encoding PI3K p85 useful for treating e.g. cancer, Type 2 diabetes, obesity -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-519374/55
                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               reagents and diagnostics.
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                                                                                                                                                                                                                                                                                                                                                                                                       2530 TTGGTAGAAGACTTGGA
                                                                                                                                                                                                                                                                                                                                                                                      17 TTGGAAGAAGACTTGAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                       15;
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                                   NISSHINBO I
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
           Kagiya T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 79;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                  (first entry)
                                     S INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.0%;
           Ichihara T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   121pp; English.
                                                                                                                                                                                                                                                                                                                      18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Murray SF, Butler MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is an antisense
                                                                                                                                                                                                                                                                                                                                                                                                             2546
                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 13.8; DB 1;
Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
           Matsumura
                                                                                                                                                                                                                                           SEQ ID NO 148
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           Moriya S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dean
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           Nishida
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
           3
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Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when

transplanting between them -

Page 121; 345pp;

Japanese.

WPI; 2002-122074/16

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Best Local
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03-JUL-2000;
10-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (ABL30512-ABL31809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas for cornea, susceptibility
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pancreas, Langerhans islet in pancreas and cornea, suscept diagnosis of genetic diseases and identifying individuals.
The present invention
                                                      Relaxin family polypeptides and antibodies recognizing them for treatment and diagnosis of metabolic disorders such as diabetes
                                                                                                   WPI; 2002-049275/06.
                                                                                                                                                                                                                                                        20-APR-2001; 2001WO-JP03399
                                                                                                                                                                                                                                                                                       01-NOV-2001.
                                                                                                                                                                                                                                                                                                                WO200181562-A1
                                                                                                                                                                                                                                                                                                                                              Unidentified
                                                                                                                                                                                                                                                                                                                                                                                      antiarteriosclerotic; immunosuppressive; cytostatic; fibroidosis; metabolic regulation; metabolic disorder; reproductive function; neurological disorder; immune disorder; scleroderma; angiogenesis
                                                                                                                                                                                                                                                                                                                                                                                                                                 Relaxin; IGF; insulin; antidiabetic; vasotropic; antiinfertility;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Relaxin/IGF/insulin family proteins related PCR primer SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABA03503;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABA03503 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18
                                                                                                                                                                                                                                                                                                                                                                        primer; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                         TAKEDA CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TTACCCTGCCACAGGAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TTACCCTGACACAAGAA 2635
                                                                                                                             Suzuki N,
                                                                                                                                                                                  ; 2000JP-0126340.
; 2000JP-0205587.
; 2000JP-0247962.
; 2000JP-0395050.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; 3 C; 6
                                                                                                                             Nishi K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%;
                            185pp; Japanese
 relates to polypeptides belonging to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ΒP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.8; DB Pred. No. 1.9e+(
                                                                                                                             Kizawa H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
.9e+02;
                                                                                                                               Harada
                                                                                                                               Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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  the
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relaxin/IGF/insulin family and their amides, esters and salts. These play a key role in metabolic regulation, especially of usage of energy sources such as sugars and lipids. The sequences can be used to prevent, treat and diagnose metabolic disorders, including disorders of the growth, proliferation and differentiation of tissues, functional lowering of reproductive functions, abnormalities of the formation of connective tissue, fibroidosis of lung, kidney or other organs, obstruction of blood circulation and internal secretions, abnormalities of body fluid balance, neurological disorders, immune disorders, scleroderma and suppression of
                                              neurological disorders, immune disorders, scleroderma and sugangiogenesis. The present sequence is a PCR primer described exemplification of the invention.
BP; 1 A;
  3 C;
     9
  <u>ଜ</u>
     ហ
     Η,
     0
     other;
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뭉
             á
                             Matches
                                   Query Match
Best Local
              1653 GGCAGGGGTCTCCGAGT 1669
                           15;
N
                                   Similarity
                             Conservative
                                  1.0%;
18
                             0;
                                   Score 13.8;
Pred. No. 1
                             Mismatches
                                    .9e+02;
                                            DВ
                                          Length 18;
                             Indels
                             0;
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Sequence

18

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Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABZ23505
                                                             22-MAY-2001; 2001EP-0201936
                                                                                                      22-MAY-2002; 2002WO-NL00322.
                                                                                                                                                28-NOV-2002
                                                                                                                                                                                          WO200295071-A2
                                                                                                                                                                                                                                                                         Replication of PCR; primer;
                                                                                                                                                                                                                                                                                                                                      Primer for analysing microsatellite repeat instability in msh-6 worms.
                                                                                                                                                                                                                                                                                                                                                                                  07-APR-2003
KONINK NEDERLANDSE AKAD VAN WETENSCHAPPE. TIJSTERMAN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA; 18
                                                                                                                                                                                                                                                                                            error; drug development; microsatellite
                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                             88.
                                                                                                                                                                                                                                                                                                   instability; msh-6;
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RESULT 201
ABZ33505/G
ID ABZ33
XX ABZ231
XX ABZ231
XX Primes
XX Replii
XX Caeno
XX PR 22-MA
XX CAENO
XX PF 22-MA
XX PF 22-MA
XX (NEVW
PA (TIJS,
XX PF 15r tl
XX PF F T plast
XX WPI;
XX PI Serve
PT replii
PT replii
PT replii
CC The sp
CC The sp The specification describes a method for determining whether a product of a gene is involved in preventing a replication error in a cell. The method comprises providing the cell with a specific inhibitor for the product and determining the level of functional expression of a marker gene in the cell, where the level of expression of the marker gene is dependent on the occurrence of the replication error. The method is used for determining whether a product of a gene is involved in preventing a replication error in a cell. The identified genes are useful for developing diagnostic tools, or as targets for drug development to manipulate cells on the basis of the presence or absence of function of the gene. Primers ABZ23504-15 were used for PCR and sequencing to analyse microsatellite instability in mesh-6 worms, in the course of the invention. Determining whether a product of a gene is involved in preventing a replication error in a cell comprises providing a specific inhibitor for the product and determining the level of expression of a marker Example 1; Page 19; 47pp; English or absence

WPI; 2003-129440/12

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RESULT 202
AAX31178/c
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                                               Query Match
Best Local S
Matches 14
                                                                                                                  AAX30947-31815 represent tag sequences of transcripts that are differentially expressed in colorectal cancer, in pancreatic cancer, or in both. The tag sequences can be used to identify genes by matching the tag to a gen data base member, or by using the tag sequences can also be used in a method colon ilbraries. The tag sequences can also be used in a method for diagnosing colon or pancreatic cancer in a sample suspected of being neoplastic. The method comprises comparing the level of at least one transcript in a first sample of a tissue to a second sample, where the first sample is a colonic tissue suspected of being neoplastic and the second sample is a normal human colonic tissue. The transcript is identified by a tag selected from AAX30947-31815. The methods of the invention can be used in the colonosis, prognosis and treatment of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 15; Conser
                                                                                                                                                                                                                                                                                                                                           Use of isolated gene transcripts - useful for developing products for the diagnosis, prognosis and treatment of cancers, particularly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tag sequence; colorectal cancer;
diagnosis; prognosis; treatment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                         Claim 2; Page 34; 120pp; English.
                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                    colon and
                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-070161/06.
                                                                                                                                                                                                                                                                                                                                                                                                           Kinzler
                                                                                                                                                                                                                                                                                                                                                                                                                                   VIND (OCYU)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-NOV-1998.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tag sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX31178;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1987 CTCCGAGAATACCTCCG
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                                                           Similarity
                        TCCAGCTCCTCCATG 1587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTCCGTGAACACCTCCG
 TCCAGCTGCTCCATG 1
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                                                                                                                                                                                                                                                                                                                                 iagnosis, prognosis and treatment pancreatic cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%;
ilarity 88.2%;
Conservative
                                                Conservative
                                                                                               BP; 4
                                                                                                                                                                                                                                                                                                                                                                                                          Vogelstein
                                                                                                                                                                                                                                                                                                                                                                                                                                   JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of a transcript increased in colorectal cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                               A;
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                                                          1.0%;
                                                                                                 3 C; 6
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Pred. No. 1.9e
0; Mismatches
                                                           Score 13.4; DB 1;
Pred. No. 1.7e+02;
                                                                                                 N
                                                Mismatches
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ss.
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                                             1.7e+02;
ches 1;
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ches 2;
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                                                                      Length 15;
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                                                Indels
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                                             Gaps
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RESULT 204
AAF51572
ID AAF515
XX
AC AAF515
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AAZ64164/c
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                                                                                                                                                                                                        Matches
                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                         The HCV sequence was screened for optimal ribozyme target sites using a computer folding algorithm and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised to target these sites and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes of the invention inhibit gene expression and/or viral replication, and are used to treat diseases associated with Hepatitis C virus (HCV) infection, e.g. cirrhosis, liver failure and hepatocellular carcinoma. The ribozymes may be used in combination with interferon to treat HCV infection, other infectious diseases, autoimmune diseases, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-APR-1998;
18-SEP-1998;
25-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Enzymatic nucleic acid; hammerhead ribozyme; virus replication; cirrhosis; liver failure; hepatocellular carcinoma; interferon; autoimmune disease; ss.
                             AAF51572
                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents the preferred target sequence of an enzymatic nucleic acid, especially a hammerhead ribozyme, which cle the Hepatitis C virus (HCV) RNA sequence at the base position given
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 83; 123pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel ribozymes for the treatment hepatitis C infection -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-062023/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Substrate for hammerhead ribozyme which cleaves HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ64164;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                               in the descriptor line
                                                                                                                    2433 CAGAATGGATAAGCC 2447
                                                                                               15
                                                                                                                                                 l Similarity
                                                                                               CAGAGTGGATAAGCC 1
                             standard;
                                                                                                                                                                                                      15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          McSwiggen JA,
                                                                                                                                                                                                      B₽;
                                                                                                                                                  Conservative
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98US-0100842.
99US-0257608.
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                                                                                                                                                              1.0%;
                             DNA;
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                                                                                                                                                 Score 13.4; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of
                                                                                                                                                                                                      0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diseases and
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                                                                                                                                                                            Length 15;
                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      conditions related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNA at
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cancer;
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30-MAR-2001 AAF51572;

(first entry)

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RESULT 205
AAF51573
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Best Local S
Matches 14
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                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45133-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other threatless.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; kelo skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasi IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris growth factor mediated cell proliferation; ichthyosis; serborrhoea; ru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell
Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic;
                                  IGF-I oligonucleotide #2533.
                                                                                                      AAF51573
                                                                                                                                    AAF51573 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                       Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                        blood vessels or any other hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              proliferation and/or inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MURD-)
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                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8; Page 77; 201pp; English.
                                                                                                                                                                                                                                                       AGGAGCAGATGACCT
                                                                                                                                                                                                                        AGGAGCAGATGACAT
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                                                                                                                                                                                                                                                                                        Conservative
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                                                                    (first
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                                                                    entry)
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                                                                                                                                                                                                                                                                                                      1.0%;
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                                                                                                                                                                                                                                                                                                                                                       G; 2
                                                                                                                                                                                                                                                                                                      Score 13.4; DB Pred. No. 1.7e+(
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                                                                                                                                                                                                                                                                                                                      Length 15;
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Matches
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Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloskin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasi IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris growth factor mediated cell proliferation; ichthyosis; serborrhoea; ru
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                                                                                                                                                        IGF-I oligonucleotide #2534.
                                                                                                                                                                                                            30-MAR-2001
                                                                                                                                                                                                                                                                                                             AAF51574 standard; DNA;
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93.3%;
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RESULT 207
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          Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
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hyperneovascular condition; hyperplasia; kidney disease;
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Pred. No. 1.7e+02;
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RESULT 208
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                                                                                                                Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; keloid; cytostatic; dermatological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease;
                                                                                                                                                                                                                                                                              30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of paoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, psoriasis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of
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                                                                                                    hyperneovascular
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                                                                                   neovascular
                                                                                                                                                                                                                                            IGF-I oligonucleotide #3626.
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                                                                                   condition; hyperplasia; ition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  any other
                                                                                                                                                                                                                                                                                                                                                  DNA;
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Pred. No. 1.7e+02;
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                                                                                                  kidney disease;
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ABX01217/c
ID ABX01217 standard; RNA; 15
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                             23-MAR-1999;
                                                                                            23-MAR-1999;
                                                                                                                                                                                                                          US2002082225-A1.
                                                                                                                                                                                                                                                                                  Hepatitis C virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MURD-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-DEC-2000
                                                                                                                                                                                                                                                                                                                                interferon gamma; consensus interrerun; перачича
substrate; hammerhead ribozyme; HH ribozyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACGTCTGGTCCTTCG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Werther GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000WO-AU00693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0140345.
                             99US-0274553
                                                                                            99US-0274553.
                                                                                                                                                                                                                                                                                                                                                                               consensus interferon; hepatotropic; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 13.4; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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RESULT 210
ABK32132/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                enzymatic nucleic acid or ribozyme is in a hammerhead (HH) or chairpin (HP) motif where the binding arms comprise sequences complementary to one of the substrate sequences defined in the specification. The HCV ribozymes are useful for modulating the expression and/or replication of HCV. They can be used to treat cirrhosis, liver failure and/or hepaticellular carcinoma. The HCV ribozymes are also useful for treating a condition associated with HCV infection in conjunction with one or more other drug therapies, particularly type I interferon, especially interferon alpha, beta or gamma or consensus interferon. The present sequence represents a substrate for a HCV hammerhead (HH) ribozyme.

Note: Some of the sequence data for this patent did not form part of the printed specification. The complete sequence data for this patent was obtained in electronic format directly from the USPTO web site
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local :
                                                                                                                                                                                                                 Human;
serial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New ribozymes targeting RNA derived from hepatitis C virus inhibit viral replication and are useful to treat hepatitis C virus infect and cirrhosis, liver failure or hepatocellular carcinoma
                                                                                                                                                                                                                                                                                                                                      ABK32132 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BLAT/)
(MCSW/)
(ROBE/)
(PAVC/)
                                                                                                                                               US6333152-B1
                                                                                                                                                                                                                                                                                                             ABK32132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 BP;
             Vogelstein B,
                                                                 20-MAY-1998;
                                                                                                                      25-DEC-2001.
                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                    cancer marker; ss.
                                                                                                                                                                                                                                                       Human colon cancer SAGE tag #233.
                                                                                                                                                                                                                                                                                  23-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          at seqdata.uspto.gov/psipsDIDEntry.html.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to enzymatic nucleic acids which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 50; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-617759/66
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                                        VIND ( OFAD)
                                                                                          20-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 2433 CAGAATGGATAAGCC 2447
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           14;
                                                                                                                                                                                                              colon cancer; colorectal cancer; pancreatic cancer; SAGE tag; analysis of gene expression; diagnostic; prognostic; probe;
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ROBERTS B.
PAVCO P A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                         CAGAGTGGATAAGCC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                        JOHNS HOPKINS
                                                                                                                                                                                                                                                                                  (first entry)
             Kinzler KW,
                                                                 98US-0081646
                                                                                             98US-0081646
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93.3%;
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            Zhang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 13.4;
Pred. No. 1.
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             zhou
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
             Ξ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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2002-153821/20

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RESULT 211
AAV48733/c
ID AAV487
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XX AAV487
XX IS-OCT
XX ErbB-2
XX Claim
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Best Local
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AAV48709-886 represent antisense oligonucleotides directed against the ErbB-2 gene. Of these, only oligonucleotides AAV48709-91 resulted in significant redcution in ErbB-2 protein expression, while oligonucleotides AAV48792-886 had little effect. The oligonucleotides exemplify the invention. The specification describes oligonucleotides that contain 8-30 nucleotides, which contain at most 8 nucleotides that can each form three hydrogen bonds to cytosine; do not contain four consecutive nucleotides able to form three H-bonds each to four consecutive cytosines; do not contain two sequences of three consecutive nucleotides able to form three H-bonds to three consecutive nucleotides able to form three H-bonds to three consecutive nucleotides able to form three H-bonds to three consecutive
                                                                                                                                                                                                                                                                                                                                                                                     Preparation of antisense oligo:nucleotide(s) which lack long runs consecutive guanosine or inosine - and have specific ratio of residues able to form two or three hydrogen bonds, have greater activity and reduced toxicity, used therapeutically or to modulate growth of cells in culture
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-JAN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ErbB-2; antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ErbB-2 gene antisense oligonucleotide ErbB-2-25.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1573
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                                                                                                                                                                                                                                                                                                                         Fig 6a;
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                                                                                                                                                                                                                                                                                                                            286pp; English.
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93.3%;
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Pred. No. 1.7e+02;
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Matches 14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-MAY-1995;
04-MAY-1995;
07-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-DEC-1994;
17-FEB-1995;
20-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytosines, and the ratio between residues able to form two H-bonds each (2R) or three such bonds (3R) is given by 2R/3R = 0.33-0.72. The oligonucleotides are used to modulate expression of genes, particularly the genes for p53, ErB-2, junB, junD, TGF-beta 1 or beta 2 to control proliferation of primary cell cultures (e.g. bone marrow stem, liver or kidney cells, osteoclasts, osteoblasts and/or keratinocytes). The oligonuclootides can also be used to analyse function of proteins (by altering their expression or activity) and therapeutically, e.g. in cases of cancer or (targeting TGF) for stimulating the immune system.
                                                                                                                                                                                                                                                                                           Beigelman L, Karpeisky A, I
Matulic-Adamic J, Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Arthritic condition; graft tolerance; immune response; target; cleavage; hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                      Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-DEC-1994;
23-DEC-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oryctolagus cuniculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX63801 standard, RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rabbit stromelysin hammerhead target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2320 CAGAGTGATGTCTGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CAGAGTGATGTGTGG
                                                                                                                                                                                                                                                                                                                                                      Gustofson J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95US-0434509.
95US-0000951.
95US-0000974.
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95US-0432874.
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                                                                                                                                                                                                                                                                                              McSwiggen J, Pavco F
A, Modak A, Usman N,
S T, Thompson JD, Wi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2334
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Pred. No. 1.9e+02;
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                                                                                                                                                                                                                                                                                   Pavco P, Stime
Usman N, Burgin A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NO:433
                                                                                                                                                                                                                                                                                                                                                      Stinchcomb
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The present invention describes a novel enzymatic having a hammerhead motif (HM) comprising: (i) at

nucleic least 5

acid (ENA) ribose

307pp;

English

Enzymatic nucleic acid molecules having a for the treatment of arthritis, induction treatment of auto-immune diseases

hammer-head motif - of graft tolerance o

WPI; 1996-300653/30.

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RESULT 213
AAT12596
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 12; Conserv
The present sequence is that of a PCR primer used for isolating the 3'-end of a cDNA sequence coding for the human protease designated T which is related to the interleukin-1 beta converting enzyme (ICB) and which induces apoptosis. First, a 600 bp fragment of the coding sequence was isolated from human monocyte cDNA using primers based on the sequence of a human genomic fragment which was 92% identical the exon 6 sequence of the human ICE gene. Then, the 5'- and 3'-ends were amplified using the Rapid Amplification of cDNA Ends (RACE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       induction; inflammation; autoimmune disease; neurodegeneration;
cancer; infection; treatment; Tx protein; anchored PCR;
polymerase chain reaction; amplification primer; ss.
                                                                                                                                                                                                                                                                         Example 1; Page 17; 88pp; French.
                                                                                                                                                                                                                                                                                                                                maturation of interleukin autoimmune diseases
                                                                                                                                                                                                                                                                                                                                                                                          New DNA encoding human
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1996-129403/13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-AUG-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ROUS ) ROUSSEL-UCLAF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Interleukin-1 beta converting enzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1586 TGAACTCCAACACCC 1600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
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                                                                                                                                                                                                                                                                                                                                                           protease(s) that induce apoptosis
kin converting enzyme, useful e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 13.4; DB
Pred. No. 2e+02;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TXA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lalanne
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ĮĻ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Livingston
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    and cause 
in treating

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RESULT 215 AAX71605 ID AAX716

AAX71605 standard; RNA;

17

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16

GTGATTACAGCTTCC

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RESULT 214
AAT35236/c
ID AAT35236/c
ID AAT35236/c
AC AAT352
XX AAT352
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XX WO9626
XX Cynthe
XX WO9626
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XX WPI
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                                                         Matches
                                                                      Query Match
Best Local Similarity
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                                                                                                                                                                                                        A series of primers (AAT35236-49) were used to sequence natural killer lytic associated protein clones (see also AAT35233) in pBluescript. Confirmation of the sequence of the 5' G-C rich region of the NKLAP gene was performed subsequent to subcloning of the clones into M13 phage. A single open reading frame coding for a 587-amino acid protein (AAR99256) was revealed.
                                                                                                                                                         Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Natural killer lytic associated protein gene primer 194.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           New human natural killer lytic associated protein -treating cancer, viral infections, graft vs. host or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Natural killer lytic associated protein; NKI cytotoxic T-lymphocyte; CTL; natural killer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT35236;
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                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 13; 65pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                         diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-412588/41
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       1375
                                                         14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
14; Conserv
GAGATTACAGCTTCC 1389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AACTGTACATGATGA 2479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AACTGTGCATGATGA 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The Tx protease converts the p30 precursor of ICE int 10 kD fragments and can be used for treating diseases ICE, e.g. inflammation. The ability to induce apoptos
                                                                                                                                                            BP; 6 A; 3 C; 4 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 6 A;
                                                         Conservative
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                                                                                93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93.3%;
                                                                                                        1.0%;
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                                                         <u>,,</u>
                                                                                Score 13.4;
Pred. No. 26
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2e+02;
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RESULT 216
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Best Local :
tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain contricion.
                                                                                                                                                                                                                                                                                                                                                                                        the Synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGP). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. ANX67275 to NAX75552 represent specific examples
                                              Vascular endothelial growth factor receptor; VEGF receptor; flt-1;
                                                                             Human KDR
                                                                                                         28-JUL-1999
                                                                                                                                                               AAX71606 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 115; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1997-259017/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Escobedo J, McSwiggen J,
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(RIBO-)
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26-OCT-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; figs-like tyroine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human KDR VEGF
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                                                                                                                                                                                                                                                                                                                                                                              nucleic acid molecules from the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present invention describes nucleic acid molecules which modulate
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                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                               GAACTTTTTAAGCTG 2418
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                                                                                                                                                                                                                                      GAACUUUUAAAGCUG 17
                                                                             VEGF
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                                                                          receptor hammerhead ribozyme substrate #618
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95US-0005974.
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Pred. No. 2e+02;
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RESULT 217
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                          Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antiporiatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                               myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome;
                                                                                                                                                                                                                               Integrin alpha 6 subunit substrate sequence SEQ ID NO:4477
                                                                                                                                                                                                                                                             19-JUN-2000
                                                                                                                                                                                                                                                                                                                     AAA21251 standard; RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 115; 218pp; English
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26-OCT-1995;
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9; Conser
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95US-0005974
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.0%;
                                                                                    syndrome; Osler-Weber-Rendu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 CC gene, an integrin alpha 6 subunit gene, or a Tie-2 gene AAA17675 to AAA17661 represent ribozyme sequences for ARNT, CC and AAA17561 to AAA17623 to AAA17684 represent their CC and AAA17167 to and AAA17560 and AAA17685 to AAA18385 and AAA19087 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA19087 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA19087 to CC AAA19154 represent their corresponding target sequences; CC and AAA19155 to AAA19222 represent their corresponding target sequences; CC AAA19231 to AAA19351 and AAA21501 to AAA21595 represent ribozyme corresponding target sequences; CC AAA19256 to AAA1688 represent their corresponding target sequences; CC AAA21689 to AAA21689 represent their corresponding target sequences; CC AAA21689 to AAA22475 and AAA23263 to AAA23342 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA23433 to CAA23432 represent their corresponding target sequences. The ribozyme of the invention are used for modulating target sequences. The ribozyme of stability of an mRNA encoding angiogenic factor, especially ARNT, critegrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are CC especially used to treat cancer, diabetic retinopathy, age related CC macular degeneration (ARMD), inflammation, and arthritis, as well as convoyabcular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, and other syndrome, and diseases related to the levels of ARNT, Tie-2, integrin cubunit alpha-6 for ARNT, Tie-2, integrin cubunit alpha-6 for particular experiences and diseases related to the levels of ARNT, Tie-2, integrin cubunit alpha-6 for ARNT, Tie-2, integrin cubunit alpha-6
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Best Local
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11-APR-2000; 2000WO-US09721.
                                               19-OCT-2000
                                                                                           WO200061729-A2
                                                                                                                                        Homo sapiens
                                                                                                                                                                                                         Ribozyme; erythropoietin; granulocyte colony stimulating factor;
                                                                                                                                                                                                                                                      Hammerhead ribozyme substrate #2471.
                                                                                                                                                                                                                                                                                                      16-FEB-2001
                                                                                                                                                                                                                                                                                                                                                    AAF04955;
                                                                                                                                                                                                                                                                                                                                                                                          AAF04955 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 8 A; 5 C; 1 G; 3 U; 0 other;
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                                                                                                                                                                                      interferon alpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              integrin subunit alpha-6, or integrin subunit beta-3.
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                                                                                                                                                                                                                                                                                                                                                                                               ВP
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Best Local Similarity
                                                                                                                                                                        Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF04956;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 2 A; 6 C; 4 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EARJ/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement
                                                                                                                                                                                                                                                                                          WPI; 2000-647423/62.
                                                                                                                                                                                                                                                                                                                                                   Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-APR-2000; 2000WO-US09721
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hammerhead ribozyme substrate #2472.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF04956 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein and interferon alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 4; Page 112; 164pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           interferon alpha;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1945 GGGCCTCTCTATGTC 1959
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                                                                                                                                                                                                                                                                                                                                                   Zwick M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zwick M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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93.3%;
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Pred. No. 2e+02;
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The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA

Claim 4; Page 112; 164pp; English

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RESULT 220
ABK02375/c
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                                                                                                                                                                                                                                                                                                                       11-FEB-2000;
28-FEB-2000;
06-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic, neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NGO0; hammerhead ribozyme; MAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; b-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CYA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkinson's disease; ataxia; Huntington's disease; reurodegenerative disease.
                                                                                                                    Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
   The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGG).

The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a
                                                                                                                                                                                                                                                               (RIBO-)
(BLAT/)
(MCSW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of crythropoietin, granulocyte colony stimulating factor protein and interferon alpha.
                                                                             Claim
                                                                                                                                                                                     WPI;
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                                                                                                        central nervous system injury
                                                                                                                                                                                    2001-607195/69
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                                                                                                                                                                                                                                                                            RIBOZYME PHARM
BLATT L.
                                                                                                                                                                                                                                                CHOWRIRA B M.
                                                                                                                                                                                                                                                              MCSWIGGEN J
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                                                                          Page
                                                                                                                                                                                                                McSwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Amberzyme
                                                                                                                                                                                                                                                                                                                       ; 2000US-181797P.
; 2000US-185516P.
; 2000US-187128P.
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                                                                          131; 200pp; English
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Pred. No. 2e
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2e+02;
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(e.g. a ribozyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 17;
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RESULT 221
ABK02376/c
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CC impartingon's disease, dementia, multiple sclerosis (MS), CC chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (MS), CC chemotherapy-induced neuropathy, and/or other neurodegenerative disease catates which respond to the modulation of NGGO expression. The CC characterist contacted with the modulat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                         Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NNGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; he-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunocytom; HIV; associated NHL; mantle-cell lymphoma; NCL; immunocytom; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; reurodegenerative disease.
                                                                                                                                                                                Synthetic
                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human NOGO
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11-FEB-2000; 2000US-181797P 09-FEB-2001; 2001WO-US04273. WO200159103-A2

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RESULT 222
ABN08362/c
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                                                                                                                                                                                                                                                                                                                                                                                       CC The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CDNAzyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a CDNAzyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a CDNAzyme) an NCH motif), a G-Cleaver (cleaving RNA with a NYN control of the presence of a divalent cation that is used CC (cleaving RNA with a YGY motif). The CD20-targetting nucleic acid is used CC cleave RNA of CD20 in the presence of a divalent cation that is CC preferably Mg.2°+. Furthermore, it may be contacted with a cell to reduce CC CD20 activity of the cell and treat a patient having a condition CC associated with the level of CD20. The treatment may further comprise the CC use of one or more therapies. In particular, the CD20 targetting CC lymphoma, low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky CC low-grade or follicular NHL, lymphocytic leukaemia, HIV (human immunocytoma (IMC), small B-cell lymphocytic lymphoma, limmune (CC immunocytoma (IMC), small B-cell lymphocytic lymphoma, immune (CC divalent cation that is preferably Mg.2°+. Furthermore, the nucleic acid cation that is preferably Mg.2°+. Furthermore, the nucleic acid cataement may further comprise the use of one or more therapies. CC treat a patient having a condition associated with the level of NOGO. The treat mervous system (CNS) injury and cerebrovascular accident (CVA, Central nervous system (CNS) injury and cerebrovascular accident (CVA, Central nervous system (CNS) injury and cerebrovascular accident (CVA, Central tectual dystrophy, and/or other neurodegenerative disease CC states which respond to the modulation of NOGO expression. The CC charkinson's disease, ataxia, Huntington's disease, Creutzfaldt-Jakob CC states which respond to the modulation of NOGO expression. The
                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-)
(BLAT/)
(MCSW/)
                Human GDMLP-1 17-mer scanning
                                                          29-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NDGO).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 88; Page 131; 200pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and central nervous system injury
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-607195/69
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-FEB-2000;
06-MAR-2000;
                                                                                                                                                                                                                                                            1573 TCCAGCTCCTCCATG 1587
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CHOWRIRA B M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RIBOZYME PHARM INC
BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                           sequence is an amberzyme molecule of the
                                                                                                                                                                                                                                  TCCAGCTCCTCCAGG 2
                                                                                                                                  standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                     BP; 4 A; 3 C; 8
                                                                                                                                                                                                                                                                                                             Conservative
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2000US-187128P.
                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                       G; 2
                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                           Score 13.4; DB 1;
Pred. No. 2e+02;
0; Mismatches 1;
                SEQ ID NO:5 sequence
                                                                                                                                                                                                                                                                                                                                                                                       U; 0 other;
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                                                                                                                                                                                                                                                                                                           1;
                                                                                                                                                                                                                                                                                                                                               Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                             invention.
                                                                                                                                                                                                                                                                                                             Indels
                  SEQ
                    ID NO:8354
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1573

TCCAGCTCCTCCATG 1587

Matches Query Match

Local

Similarity

93.3%;

Score 13.4; Pred. No. 26 Mismatches

4; DB 1; 2e+02;

Length 17;

<u>,</u>

Gaps

0

Conservative

0

Sequence 17

BP; 6 A; 2 C; 8 G; 1 T; 0 other;

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of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific hiomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1
                                                           N.B. The sequence data
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 substrates, of hGDMLP-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein 1 (hGDMLP-1). The protein and polymeration regions hGDMLP-1 can be used in gene therapy and vaccine production.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
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30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (AEOM-) AEOMICA INC.
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30-JAN-2001;
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                                                                                         .nvention.
ftp.wipo.int/pub/published_pct_sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (hGDMLP-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention describes a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID 8354 214pp; English
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2001WO-US00665.
2001WO-US00666.
2001WO-US00667.
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                                was obtained in electronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       describes a human genome-derived myosin-like The protein and polynucleotide sequences of
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TCCAGCTCCTCCTTG

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RESULT 223
ABN08363/c
           The present invention describes a human genome-derived myosin-like CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of CC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 nucleic acids can be used as probes to detect, characterise CC and quantify hGDMLP-1 nucleic acids in samples, as amplification CC substrates, to provide initial substrates for the recombinant engineering CC of hGDMLP-1 protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific concentration and/or amount specifically of hGDMLP proteins, as specific CC deficiency in hGDMLP-1 production, and in vaccines or for replacement CC diagnosing a disorder associated with the expression of hGDMLP-1; in CC diagnosing a disorder associated with the expression of hGDMLP-1; in localised to chromosome 22. The present sequence represents an oligomer used in the present screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                         New polypeptide, for raising antibodies that proteins, or as specific biomolecule capture surface-enhanced laser desorption ionization,
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30-JAN-2001;
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27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                              myosin-like
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AEOMICA INC.
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2001WG-US00661.
2001WG-US00662.
2001WG-US00663.
2001WG-US00665.
2001WG-US00665.
2001WG-US00665.
2001WG-US00666.
2001WG-US00667.
2001WG-US00667.
2001WG-US00667.
2001WG-US00669.
2001WG-US00669.
2001WG-US00669.
                                                                                                                                                                                                                                                                                                                                              protein hGDMLP-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-mer scanning
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                                                                                                                                                                                                                                                                                                                                                                        recognize hGDMLP-1 probes for
                                                                                                                                                                                                                                                                                                                                                            comprises human
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RESULT 224
ABNO83
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ABNO84
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ABNO86
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ABNO8
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30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                         New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1 -
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N.B. The sequence data for this patent did not form precification, but was obtained in electronic format at ftp.wipo.int/pub/published_pct_sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                                                                                        WPI; 2002-179446/23.
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                                                                                                                                                     SEQ ID 8356; 214pp; English
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2001WO-US00665.
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2001WO-US00667.
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2001US-266860P
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2000US-236359P.
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protein 1 (hGDMLP-1). The protein and polynuclectide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, character and quantify hGDMLP-1 nucleic acids in samples, as amplification

invention describes a human genome-derived myosin-like

The present

substrates, to provide initial substrates

the recombinant engineering

detect, characterise as amplification

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RESULT 225
ABN08381/c
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Best Local
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                                                                                                                                                            30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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            WPI; 2002-179446/23
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27-SEP-2000;
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Similarity 93.3%;
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Query Match

Local

1.0%;

Score 13.4; DB Pred. No. 2e+02;

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Length

17;

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substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific polypeptides or surface-enhanced laser desorption instation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present revent
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Sequence 17 BP; 5 A; 5 C; 5 G; 2 T; 0 other;
                                                                                N.B. The sequence data for this patent did not specification, but was obtained in electronic i
                                                                                                                                            invention.
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ARBSULT 226
ARBNO8384/c
ID ARBNO83
XX ABNO83
XX ABNO83
XX PAMAY
DE Human;
KW Human;
KW Muscle
XX Skelet
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX PR 25-MAY
XX Z6-MAY
PR 27-SEP
PR 27-SEP
PR 27-SEP
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PR 27-SEP
PR 30-JAN
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PR 30-JAN 밁 Ś Matches 26-MAY-2000; 21-SEP-2000; 27-SEP-2000; 04-OCT-2000; Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 30-JAN-2001; 25-MAY-2001; WO200192524-A2 Homo sapiens. 29-MAY-2002 (first entry) ABN08384 standard; DNA; 17 Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:8376. 1507 CAGCCGGCTGTGCAC 1521 17 1 Similarity
14; Conserv CAGCTGGCTGTGCAC 2001WO-US00661. 2001WO-US00662. 2001WO-US00663. 2001WO-US00664. Conservative 2001WO-US16981 2000GB-0024263. 2000US-234687P. 2000US-236359P. 2000US-207456P. 0; Mismatches ۲. Indels 0 Gaps

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              WO200192524-A2
                                          Homo sapiens.
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                                          ABN08952
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ABN08952;

ABN08952 standard; DNA; 17

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                                                          Query Match
Best Local S
Matches 14
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                                                                                                                          Sequence 17 BP; 4 A; 4 C; 8 G; 1 T; 0 other;
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                                                                                                                                                                          specification,
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27-SEP-2000;
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                                                                                                                                                                                                           invention
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                                                                             Local Similarity
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                              CAGCGCCTGGAAGAG 1369
                                                              Conservative
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2001WO-US00670
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2000US-236359P
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Pred. No. 2e+02;
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Human GDMLP-1 17-mer scanning

SEQ ID NO:5

sequence SEQ

hGDMLP-1; heart; ID NO:8944.

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genome-derived myosin-like protein 1;

29-MAY-2002

(first entry)

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The present invention describes a human genome-derived myosin-like CC protein 1 (hGpMLP-1). The protein and polynucleotide sequences of CC hGpMLP-1 can be used in gene therapy and vaccine production. The CC hGpMLP-1 nucleic acids can be used as probes to detect, characterise CC hGpMLP-1 nucleic acids can be used as probes to detect, characterise CC substrates, to provide initial substrates for the recombinant engineering CC of hGpMLP-1 protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGpMLP-1 proteins or polypeptides may CC be used as immunogens to raise antibodies that specifically recognise CC hGpMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGpMLP proteins, as specific CC deficiency in hGpMLP-1 production, and in vaccines or for replacement CC diagnosing a disorder associated with the expression of hGpMLP-1, in CC diagnosing a disorder associated with the expression of hGpMLP-1, in CC chromosome 22. The present sequence represents an oligomer used in the greening of the hGpMLP-1 sequence in the exemplification of the present
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21-SEP-2000;
27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human
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                                           N.B. The sequence data for this patent did not specification, but was obtained in electronic i
                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID 8944; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                        myosin-like protein hGDMLP-1
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                           ftp.wipo.int/pub/published_pct_sequence.
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therapy; vaccine; heart disease;
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Query Match

Sequence 17

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A; 5 C; 8 1.0%;

G; 1 T; 0 other; Score 13.4; DB

1;

Length 17;

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RESULT 229
ARNO89
XX ABNO89
XX ABNO89
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XX Human;
XX Human;
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XX WO2001
XX WO2001
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                              of hGDMLP-1 protein variants having desired phenotypic improvements, at for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specifically of hGDMLP proteins, as specifically of hGDMLP proteins.
                                                                                                                                                                                                                The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1
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27-SEP-2000;
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                            biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific
                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure;
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in hGDMLP-1 production, and
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0; Mismatches
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therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present

screening of the hGDMLP-1 sequence invention.

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RESULT 230
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                                               New polypeptide, for raising antibodies that proteins, or as specific biomolecule capture surface-enhanced laser desorption ionization,
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                   Disclosure;
                                       myosin-like
                                                                                        WPI; 2002-179446/23.
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                                                                                                                               AEOMICA INC.
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                 SEQ ID 9000; 214pp; English
                                      protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                    2001WO-US00666.
                                                                                                                                                                                                                                                                                  2000US-207456P
2000US-234687P
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                                                                                                                                                            2001WO-US00670
                                                                                                                                                                                           2001WO-US00667
                                                                                                                                                                                                              2001WO-US00665.
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                                                                                                                                                                                                                                                     2001WO-US00661.
                                                                                                          Penn SG,
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                                       hGDMLP-1
                                                                                                          Hanzel DK,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13.4; DB Pred. No. 2e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO:5 sequence SEQ
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                                                                                                         Rank DR,
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                                                        recognize hGDMLP-1 probes for
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                                                 comprises human
                                                                                                          Chen W,
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                                                                                                          Shannon ME
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The

present invention describes a human genome-derived myosin-like

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AEOMICA

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of CC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 nucleic acids can be used as probes to detect, characterise CC and quantify hGDMLP-1 nucleic acids in samples, as amplification (CC substrates, to provide initial substrates for the recombinant engineering CC of hGDMLP-1 protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may cC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific concentration and/or amount specifically of hGDMLP proteins, as specific concentration, as therapeutic supplement in patients having specific constantion, as therapeutic supplement in patients having specific confisation, as therapeutic supplement in patients having specific confisation, and hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for cdiagnosing a disorder associated with the expression of hGDMLP-1, in CC cdiagnosing a disorder associated with the expression of hGDMLP-1, in the correspondence of the hGDMLP-1 sequence represents an oligomer used in the correspondence of the hGDMLP-1 sequence in the exemplification of the present
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N.B. The sequence data for this patent did not not necessarily and in electronic interpretation. But was obtained in electronic interpretation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine;
                                                                                                      30-JAN-2001;
30-JAN-2001;
                                                                                                                                          30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                 27-SEP-2000;
                                                                                                                                                                                                                                                                       21-SEP-2000;
                                                                                                                                                                                                                                                                                      26-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        skeletal muscle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:9003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABN09011;
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                                                                     30-JAN-2001;
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                                                                 2001WO-US00666.
2001WO-US00667.
2001WO-US00668.
2001WO-US00669.
                                                                                                                                        2001WO-US00662.
2001WO-US00663.
2001WO-US00664.
2001WO-US00665.
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                                 2001WO-US00670
2001US-266860P
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2000GB-0024263.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chromosome 22; gene therapy; vadisorder; amplicon; screening;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
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Pred. No. 2e+02;
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t directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hGDMLP-1; heart; heart disease;
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RESULT 232
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of CC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 nucleic acids can be used as probes to detect, characterise CC and quantify hGDMLP-1 nucleic acids in samples, as amplification consumering conformation variants having desired phenotypic improvements, and CC of hGDMLP-1 protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may CC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption CC ionisation, as therapeutic supplement in patients having specific CC deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for CC diagnosing a disorder associated with the expression of hGDMLP-1, in CC chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
16-JUL-2001;
20-JUL-2001;
13-NOV-2001;
19-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N.B. The sequence data for this specification, but was obtained at ftp.wipo.int/pub/published_po
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gu Y,
                                                                                                                                                                                      Eating disorder; polymorphism; dataset; allele; HGBASE identification; serotonin receptor 1D; delta-opioid receptor; dopamine receptor D2; anorexia nervosa; bulimia nervosa; probe; ss.
                                                                                                                                                                                                                                               Opioid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure;
                                                                                                                                                            Unidentified
                                                                                                                                                                                                                                                                             12-JUN-2003
                                                                                                                                                                                                                                                                                                                                    ABT34296 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17
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                                                                      16-JUL-2002; 2002WO-US22555
                                                                                                      13-FEB-2003
                                                                                                                               WO2003012143-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        edification, but was obtained in electronic ffp.wipo.int/pub/published_pct_sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                               1506 CCAGCCGGCTGTGCA 1520
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14;
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                                                                                                                                                                                                                                             receptor
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                                                                                                                                                                                                                                                                                                                                                                                                          CCAGCCGGCGGTGCA 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein hGDMLP-1
2001US-305153P.
2001US-306440P.
2001US-331285P.
2001US-340843P.
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                                                                                                                                                                                                                                             probe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               recognize hGDMLP probes for , comprises human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      directly from WIPO
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RESULT 233
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XX ABT388
XX ABT388
XX ABT388
XX Cytost
KW Cytost
KW Schizo
KW Schizo
KW Homo s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel isolated nucleic acid molecule comprising a variant gene associated with an eating disorder and selectifrom any of 119 polymorphisms with their corresponding genotyping in dataset, alleles and HGBASE identification, given in the specification. The novel nucleic acid molecule has polymorphisms in the serotonin receptor 1D, delta-optioid receptor, or dopamine receptor 2D, which is useful in diagnostic and prognostic assays for eating disorders, in particular anorexia nervosa and bulimia nervosa. This polynucleotide sequence represents a opioid receptor 1D probe of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
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                                                                                  New isolated nucleic acid, useful for treating viral disea associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells
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                                 Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                       17-SEP-2001; 2001FR-0011978
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003025175-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tumour suppression related human fukutin oligo
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                                                                                                                                                                                                                                                                                                                                                                                                                                              17-SEP-2002; 2002WO-IB04208
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                                                                                                                                                                                                                                                                                                                             MOLECULAR ENGINES LAB
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                              Page
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                                                                                                                                                                                                                                                                   Amson R,
                                 554;
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                              720pp;
                                                                                                                                                                                                                                                                      Tuijnder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 13.4; DB 1;
Pred. No. 2e+02;
                                 French
                                                                                                                                                                                                                                                                         z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disorder and selected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 234
ABT38840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cc for detecting, identifying, quantifying and/or amplifying a nucleic acid, cc e.g. as one component of a gene chip, in vitro as (antilsense reagents, cc and for production of recombinant polypeptides. Any of the nucleic acids, cc polypeptides, vectors containing the mucleic acids, cells containing the cvector or antibodies directed against the polypeptides are useful for greparation of pharmaceuticals for prevention and/or treatment of viral cliesases that are characterised by development of tumours or cell cdseases that are characterised by development of tumours or cell cdseases that are characterised by development of tumours or cell cdseases that are characterised by development of tumours or cell cdseases. Analysis of the expression of the 17 mer nucleic acids in cc schizophrenia. Analysis of the expression of the 17 mer nucleic acids in cc diseases. The polypeptides can also be used to generate antibodies, and batten temples is useful for diagnosis and/or prognosis of these components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression crelated human fukutin oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers
                                                                New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-SEP-2002; 2002WO-IB04208.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17
                                                                                                                                                                                                                    WPI; 2003-313353/30
                                                                                                                                                                                                                                                                                        Telerman A,
                                                                                                                                                                                                                                                                                                                                                                    (MOLE-) MOLECULAR ENGINES LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                        17-SEP-2001; 2001FR-0011978.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2003025175-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tumour suppression related human fukutin oligo SEQ ID No 4477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABT38840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABT38840 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1930 GCCTGCACACAGGAT 1944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCCTACACACAGGAT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 3 A; 3 C; 6 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                        Amson R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
   557;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.0%;
720pp; French
                                                                                                                                                                                                                                                                                        Tuijnder M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13.4; DB 1; Length 17; Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 20+1
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15

WPI; 2003-129306/12

(BIOI-) BIOINVEST LTD

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ABX15000/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      consecutive nucleotides from the 17 mer sequence, a sequence with, after coptimal alignment, at least 80 % identity to the 17 mer sequence, a complement of any of them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers considered nucleic acids of the invention are useful as probes and primers component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, ce.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, completely vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral components of pharmaceuticals for prevention and/or treatment of viral componentation, specifically cancer but also Alzheimer's disease and componentation, specifically cancer but also Alzheimer's disease and the sequences of the sequences of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these components of these components of these components of these components at the polypeptide and antibodies are useful as components of protein the polypeptide and antibodies are useful as components of protein the polypeptide and antibodies are useful as components of protein the polypeptide and antibodies are useful as components of protein the polypeptide and antibodies are useful as components of protein the polypeptide and antibodies are useful as components of protein the polypeptide and antibodies are useful as components of protein the polypeptide and antibodies are useful as components of protein the polypeptide components of the polypeptide components of the polypeptide components of protein the polypeptide components of the polypeptide components of protein the polypeptide components of protein the polypeptide components of protein th
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-MAR-2003 (first entry)
                                                                                                                              11-MAY-2001; 2001US-290016P
                                                                                                                                                                          13-MAY-2002; 2002WO-US14940.
                                                                                                                                                                                                                                                                   WO200292838-A2
                                                                                                                                                                                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human delta opioid receptor OPRD1-5 SNP genotyping PCR probe #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABX15000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABX15000 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 4 A; 2 C; 8 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1861 GATCTGGTGTCAGAG 1875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        delta opioid receptor; OPRD1-5; ss; PCR; probe; SNP; nucleotide polymorphism; eating disorder; anorexia n homeostasis disorder; chromosome 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GATCGGGTGTCAGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                      moiety"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                             rhodamine (TAMRA)
                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "G is covalently linked
                                                                                                                                                                                                                                                                                                                                                                           *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                             note= "T is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%;
                                                                                                                                                                                                                                                                                                                                                                             b
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                             covalently linked
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 13.4; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1;
                                                                                                                                                                                                                                                                                                                                    to a 6-carbotetramethyl-
                                                                                                                                                                                                                                                                                                                                                                                                                                             ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 anorexia nervosa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                             þ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                TET
                                                                                                                                                                                                                                                                                                                                                                                                                                             (not defined)
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RESULT 236
ABZ61761/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modulates the expression of the nucleic acid, diagnosing a genetic correction of predisposition to an eating or energy homeostasis disorder by detecting the presence or absence of the variant nucleic acid in a patient sample, can allele specific primer that detects a polymorphism in the gene encoding a delta opioid receptor associated with an eating or energy contain the variant nucleic acids. The variants are named OpRD1-1 contain the variant nucleic acids. The variants are named OpRD1-1 contain the variant opioid receptor gene is located on chromosome 1. The nucleic acid molecules and delta opioid receptor variant are useful for diagnosing a genetic predisposition to an eating or energy homeostasis disorder, such as anorexia nervosa. The allele specific primer is useful for detecting polymorphism in the gene encoding a contain the genetic opioid receptor associated with the disorder cited. The present sequence is a genotyping PCR probe for detecting the presence of a particular SNP (single nucleotide polymorphism) in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                  Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    homeostasis disorder. Also included are a delta opioid receptor variant encoded by the nucleic acid, an isolated antibody that specifically recognises the delta opioid receptor variant, a vector comprising the nucleic acid, a host cell transformed to contain the vector, producing the polypeptide by culturing the host cell, identifying an agent which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated nucleic acid molecule encoding a delta opioid receptor variant associated with an eating or energy homeostasis disorder, useful for diagnosing a genetic predisposition to such disorder, e.g.
                                                                                                                                                                                                                                                                                                                                                                      Human H-Ras DNAzyme target #552.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to an isolated nucleic acid molecule encoding delta opioid receptor variant associated with an eating or energy
                Mcswiggen
                                                                                                   29-MAY-2001;
06-JUN-2001;
                                                                                                                                                         29-MAY-2002; 2002WO-US16840
                                                                                                                                                                                            05-DEC-2002
                                                                                                                                                                                                                               WO200297114-A2
                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                             21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                              ABZ61761;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABZ61761 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example; Page 19; 39pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      anorexia nervosa
                                                 (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1831 AAAGATGATGCCACA 1845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14;
                                                   RIBOZYME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAAGATGAGGCCACA
                                                                                   ; 2001US-294140P.
; 2001US-296249P.
; 2001US-318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 1 A; 4 C; 5 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                   PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.4; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            variant
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Query Match
Best Local
                                                                                                          The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ra human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABS25989 - ABS26216, ABS24544 - ABS26531, ABS26520 - ABS26524, ABS26531 - ABS26535 represent substrate/target sequences for the human ribozymes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
                                                                     Sequence 17 BP; 3 A; 4 C; 8 G; 2 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 58; Page 121; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treating cancer, mountained HER2, K-Ras, H-Ras, N-Ras,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-140484/13.
Local
                        1.0%;
  Pred.
                        Score 13.4;
                      DB 1;
                        Length 17;
                                                                                                                                                                                                                                                                                                                                         N-Ras,
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Matches
ABZ64832 standard;
                                                                  2283 GGCTCCAGAAGCCCT 2297
                                               17
                                                                                   l Similarity
14; Conserv
                                               GGCTCCAGCAGCCCT 3
                                                                                     Conservative
 RNA; 17
                                                                                             93.3%;
 BP.
                                                                                    <u>,</u>
                                                                                   Mismatches
                                                                                             No. 2e+02;
                                                                                     1;
                                                                                     Indels
                                                                                     0
                                                                                     Gaps
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밁 S

RESULT 237
ABZ64832
ID ABZ648
XX ABZ648
XX ABZ648
XX ABZ648
XX Human
XX Human
XX Homo s
XX Homo s
XX WO2002
XX WO2002
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XX Homo s
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XX Novel
PT treati
PT treati
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CC acid in
CC acid in Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss. 05-DEC-2002. WO200297114-A2. Homo sapiens. Human HER2 DNAzyme substrate #289.

21-MAR-2003

(first entry

29-MAY-2002; 2002WO-US16840

29-MAY-2001; 06-JUN-2001; 10-SEP-2001; ; 2001US-294140P. ; 2001US-296249P. ; 2001US-318471P.

(RIBO-) RIBOZYME PHARM INC

Mcswiggen

2003-140484/13

Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences -

Claim 4; Page 138; 185pp; English.

The invention relates to a novel short interfering RI acid molecule or an enzymatic nucleic acid molecule, expression of a nucleic acid molecule encoding HER2, RNA (siRNA) nucleic e, that modulates 2, K-Ras, H-Ras, N-Ra

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ABZ65210
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Best Local :
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                                       The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ra human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ66585 represents substrate/target accommences for the human riboxymae of the invention substrate/target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ6216, ABZ64544 - ABZ6531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention.
                                                                                                                                                                                                                                                                                    Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encod HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-MAY-2001; 2001US-294140P.
06-JUN-2001; 2001US-296249P.
10-SEP-2001; 2001US-318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
                                sequences
                                                                                                                                                                                                                                                    Claim 4; Page 145; 185pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                             Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC
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                              for the human ribozymes of the invention.
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Pred. No. 2e+02;
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Sequence 17 BP; 3 A;

3 C;

7

G; 4 U; 0 other;

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RESULT 239
ABZ65221
XX ABZ652
XX ABZ652
XX ABZ652
XX Human;
AK Human;
KW Human;
KW enzyma
KW enzyma
KW Human;
KW enzyma
KW Human;
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Best Local Similarity
Matches 11; Conserv
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06-JUN-2001;
10-SEP-2001;
                                                                                                                                                                                                                                                               acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-RAS, H-RAS, N-R human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-Theumatic activity. The nucleic acid molecule activity. The nucleic acid molecules are useful for reducing HER2, K-RAS, H-RAS, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ6434 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66530 - ABZ66530 - ABZ66531 sequences for the human ribozymes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel short interfering RNA (siRNA) nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 146; 185pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
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10; Conserv
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CAGAGUGAUGUGUGG
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; 2001US-296249P.
; 2001US-318471P.
                                                                                                           Conservative
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                                                                                                                               1.0%;
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16
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                                                                                                                                  Score 13.4;
Pred. No. 26
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                                                                                                           Mismatches
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                                                                                                                                     2e+02;
                                                                                                                                                            DB 1;
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                                                                                                        0,
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RESULT 241
ABF46427/c
ID ABF464
XX ABF464
XX ABF464
XX CAFFB
XX Oligon
XX SNP, s
KW SNP, s
KW Centra
XX Centra
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.

ABC00010-ABE09989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABE09989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABE09081 for the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences.
               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         designed to detect single nucleotide polymorphisms and methylation status
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-APR-2000; 2000DE-1019173
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                                                                               Oligonucleotide SEQ ID NO 146424 for detecting
                                                                                                                 21-FEB-2002
                                                                                                                                                 ABF46427;
                                                                                                                                                                               ABF46427 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                              Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID 146423; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                             1872 AGAGATGGAGATG 1884
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                                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                                                 (first entry)
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100.0%; Pred. No. 1.6
ive 0; Mismatches
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hes 0;
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                                                                                 SNP TSC0036912.
                                                                                                                                                                                                                                                                                                                                                          Length 13;
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RESULT 242
ABF4944
XX ABF494
XX ABF494
XX Oligon
XX Oligon
XX SNP; 8
KW SNP; 8
KW Centra
XX Centra
XX Homo 8
XX Gentra
XX Homo 9
XX Gentra
XX Homo 9
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XX Gent
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ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide SEQ ID NO 149443 for
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                                                                                                         07-APR-2000; 2000DE-1019173
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RESULT 243
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    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                           Set of oligonuclectides, useful for diagnosis and cell typing, is designed to detect single nuclectide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                              designed to methylation
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genomic DNA. of cancer an

Best Local Similarity

100.0%; 0.9%;

Score 13; DB 1; L Pred. No. 1.6e+02; 3 T; 0 other;

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Query Match

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                                         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.

ABC00010-ABC99989, ABF00010-ABF99889, ABH000110-ABH99989 and ABI000110-ABC99989, Tepresent the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                          Claim 1; SEQ ID 206165; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                           Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single nucleotide polymorphisms and cytosin
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larity 100.0%;
Conservative (
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RESULT 245
ABH06189/c
ID ABH061
XX ABH061
XX ABH061
XX Oligon
XX SNP;
KW SNP;
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KW SNP;
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XX Homo s
PN WO2001
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XX WO2001
XX O1-APR
XX O7-APR
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RESULT 246
ABH51404/c
ID ABH514
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Best Local S
Matches 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single nucleotide polymorphisms and cytosine methylation status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.

ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH9989 and ABC00010-ABH82073 represent the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-APR-2001; 2001WO-IB00713.
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                               ABH51404;
                                                                                       ABH51404 standard; DNA; 13
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13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 3 A; 4 C; 0 G; 6 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                               s; Score 13; DB; Pred. No. 1.6
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                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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RESULT 247
ABH51405
ID ABH514
XX ABH514
XX ABH514
XX ABH514
XX OBH514
XX ORD;
C2-FEB
XX SNP;
CW SNP;
CW Centra
XX SNP;
CONT CANTON S
XX SNP;
CONT CANTON S
XX ORD;
CANTON S
XX ORD;
CONT CANTON S
XX O
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.

ABC00010-ABC9999, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Set of oligonucleotides, useful for diagnosis and cell designed to detect single nucleotide polymorphisms and methylation status \, -
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                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88;
                                                                                                                                                                                                                                                                                               Oligonucleotide SEQ ID NO 251382 for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 13
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                                                                                                                 Homo sapiens.
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                                                                                                                                                                           central nervous
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13; Conserv
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                                                                                                                                                                           system;
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                                                                                                                                                                           gastrointestinal;
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                                                                                                                                                                                                                                                                                               detecting
                                                                                                                                                                              respiratory; immune; metabolic
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ARESULT 248
AAF51570
ID AAF515
XX AAF515
XX AAF515
XX AAF515
XX AAF516
XX AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP3; IGFBB3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE39989, ABF00011-ABF39989, ABF00011-ABF39989, ABF00011-ABF39989, and ABI00010-ABIB2073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
   Wraight CJ,
                                                                                                                                                                                21-JUN-2000; 2000WO-AU00693
                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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larity 100.0%;
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Pred. No. 1.6e+(
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBD3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease;
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Ameliorating the effects of a disorder,

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                                                                                                                                                                                                                                                                                                                                                                   IGF binding protein; IGFBP-2; IGFBP3; inflammation; poriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
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                          Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an analysis acid that inhibits or reduces growth factor mediated
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The present invention relates to a method of skin disorders. The method comprises co

od for ameliorating the effects contacting the skin with an

Example

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antis nucleic acid that inhibits or reduces growth factor mediated cell

an antisense

and/or inflammation

201pp;

English

21-JUN-1999;

99US-0140345.

(MURD-) MURDOCH CHILDRENS RES INST

2001-041421/05

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Werther GA,

Edmondson SR,

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ARESULT 251
AAP5293/C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neovascular condition of the retina;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2333 GGTCCTTCGGGGT 2345
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13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; DNA;
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Pred. No
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Matches 13
The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an
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                                                                                                                                                                                                                                                                                                                             administering UV (ultra-violet) treatment nucleic acid that inhibits or reduces grow
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                                                                                                                                                                                                                                    Example
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Pred. No.
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                                                                                                                                                    oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see ARF45131 and ARF45131-845161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of
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Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological, cardiant, virucide; ophthalmological, keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.

WO200078341-A1

28-DEC-2000

21-JUN-2000; 2000WO-AU00693.

99US-0140345.

(MURD-) MURDOCH CHILDRENS RES INST.

WPI; 2001-041421/05.

Wraight CJ,

Werther GA,

Edmondson

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an incleic acid that inhibits or reduces growth factor mediated proliferation and/or inflammation an antisense cell

Example 8; Page 86; 201pp; English.

of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAP45151 and AAP45153-F45161). The method is useful for ameliorating the effects of AAP45153-F45161). The method is useful for ameliorating the effects of present invention relates to a method for ameliorating the effects ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids ဇ္ဌ

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  Query Match
Best Local S
Matches 13
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Best Local S
Matches 13
                                                                                                                                                                         The present invention provides the protein, cDNA and genomic sequences of human interleukin-13 (III3), and describes the single nucleotide polymorphisms (SNPs) found within the gene, which is found on chromosome 5q31. III3 is a pro-inflammatory cytokine thought to be involved in the pathogenesis of asthma and other immune and inflammatory diseases. The III3 sequences and the SNPs identified can be used in drug screening, to determine an individual's susceptibility to disease, in forensic and paternity testing, and to identify treatments for cancer, immune and inflammatory diseases, including asthma and diseases characterised by fibrosis. The present sequence is an III3 allele-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other selector disease, kidney disease, hyperproliferation of the inside of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel polynucleotide comprising single nucleotide polymorphisms human interleukin-13 gene is useful for studying expression and function of interleukin-13, as well as diagnosing and treating of the control of the con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; interleukin-13; II13; single nucleotide polymorphism; SNP; cancer; inflammation; immune disorder; cytokine; asthma; chromosome 5q31; fibrosis; forensic; disease susceptibility; drug screening; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 15; Page 20; 85pp; English.
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                                                                                                    Sequence 15
                                                                                                                                                      oligonucleotide.
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     l Similarity
13; Conserv
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13; Conserv
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ilarity 100.0%;
Conservative
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0.9%; Score 13; ilarity 100.0%; Pred. No. Conservative 0; Mismatcl
                                                                                                    BP; 4
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Pred. No.
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                                                                                                    T; 0 other;
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2e+02;
                                                    DB 1;
                             2e+02;
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1649 TGCTGGCAGGGGT 1661

Matches Query Match

Local

l Similarity

0.9%; Score 13; 100.0%; Pred. No.

DB 1; 2e+02;

Length 15; Indels

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Mismatches

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Gaps

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RESULT 255
ABK41344/c
ID ABK413
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                                                                                                                                                                       human translation initiation factor 2B gamma subunit, eIP2Bgamma, and proteasome alpha subunit 1 ,PMSA1, acting on Hepatitis C virus, HCV, sequences) and a target recognition sequence of recovered ribozymes are sequenced to identify the cellular regulator. Also included are target sequence tags, TST, derived from eIF2Bgamma and pMSA1, the ribozyme sequence tags, RST, targetting the TSTB (and a list of target genes given in the specification), methods of identifying the ribozyme sequences and other compounds having a positive or negative effect on viral replication or interaction with the cellular regulator. The methods are useful for identifying a compound that modulates the activity of a viral cellular regulator, identifying
                                a ribozyme reactive with a cellular regulator of virus replication or expression, and for treating an HCV infection by inhibiting the activit of a cellular regulator involved in HCV replication. The ribozymes and inhibitory compounds identified by the above screening methods are used to reduce the severity of such an infection. The methods allow rag propagation or pathogenesis of infectious agents. The present sequence is a ribozyme sequence tag, RST, of the invention.
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a randomised ribozyme gene vector library which is introduced into a population of cells expressing negative selection marker gene operatively linked to viral nucleic acid acte by cellular regulator of virus replication or expression (e.g. the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              eIF2Bgamma; ribozyme; ribozyme sequence tag; RST; TST; target sequence tag; HCV; hepatitis C virus infection; virucide; hepatotropic; antiinflammatory; proteasome alpha subunit; PMSA1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying cellular regulators essential in pathogenesis of infectious agents, useful for treatment of infectious diseases preferably viral diseases especially hepatitis C virus (HCV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human eIF2Bgamma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 16; Fig 4D; 74pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-MAY-2000; 2000US-0563794.
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B₽;
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4 A;
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4 C; 5 G; 2 U; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gamma subunit;
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RESULT 256
ADX63934/c
ID ADX639
XX AAX639
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XX AATChri
DE Rabbit
XX Arthri
KW Arthri
KW Hammex
KW Strome
KW Cream
XX Orycto
PN WO9618
PN W
                  The present invention describes a novel enzymatic nucleic acid (ENA) CC having a hammerhead motif (HM) comprising: (i) at least 5 ribose CC residues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) CC at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification. CT he ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, cc particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for treating graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly present invention.
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23-DEC-1994
23-DEC-1994
23-DEC-1994
17-FEB-1995
20-APR-1995
02-MAY-1995
04-MAY-1995
07-JUL-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic nucleic acid molecules having for the treatment of arthritis, induction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Draper K, Gustofson J, McS
Beigelman L, Karpeisky A, I
Matulic-Adamic J, Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Draper K, G
Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oryctolagus cuniculus.
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it of auto-
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95US-0000951.
95US-0000974.
95US-0512861.
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95US-0390850.
95US-0426124.
95US-0432874.
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94US-0363253.
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A, Modak A, Usman
s T, Thompson JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             target
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pavco P, Stinch Usman N, Burgin JD, Wincott F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hammer-head motif - used of graft tolerance or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NO:566
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RESULT 257
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Matches 13; Conserv
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23-DEC-1994;
17-FEB-1995;
20-APR-1995;
02-MAY-1995;
04-MAY-1995;
07-JUL-1995;
07-JUL-1995;
The present invention describes a novel enzymatic nucleic acid (ENA) having a hammerhead motif (ENA) comprising: (i) at least 5 ribose residues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification. The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for enhancing graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENA's can also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arthritic condition; graft tolerance; immune response; target; cle hammerhead ribozyme; hairpin ribozyme; huann; rabbit; mouse; colle stromelysin; synovial membrane; joint; arthritis; ostcoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                          Enzymatic nucleic for the treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oryctolagus cuniculus.
                                                                                                                                                                                                                                                                                                      Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matulic-Adamic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Beigelman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Draper K,
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13-DEC-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rabbit stromelysin hammerhead target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-JUL-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RIBOZYME
                                                                                                                                                                                                                                                                                                   1; Page 154; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTGCTGAAGGAAG 2428
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                                                                                                                                                                                                                                                                                                                                                      of auto-immune diseases
                                                                                                                                                                                                                                                                                                                                                                          nucleic acid molecules having a reatment of arthritis, induction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gustofson J, McSv
L, Karpeisky A, I
amic J, Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%; Scilarity 100.0%; E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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95US-0000974.
95US-0512861.
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95US-0426124.
95US-0432874.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Α,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          McSwiggen J, Pavco P, Sti
A, Modak A, Usman N, Burg
B T, Thompson JD, Wincott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  w
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  σ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pavco P, St....
Usman N, Burgin A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
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                                                                                                                                                                                                                                                                                                                                                                          hammer-head
of graft tol
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stinchcomb
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Best Local
The present invention describes an enzymatic nucleic acid molecule (I) with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDNA sequence encoding maize belta-9 desaturase. (I) can be used to modulate expression of a gene, preferably belta-9 desaturase or a granule bound starch synthase (GBSS) gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate caffeine synthesis in a coffee plant, nicotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a tobacco, aspen, poplar or pine plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Edington BI
Merlo PAO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly specific. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               granule bound starch synthase; hammerhead ribozyme; hairpin ribomodulation; gene expression; transgenic plant; cleavage; canola caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
                                                                                                                                                                                                                                                                                                                                                                      Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound synthase in maize or canola
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Maize; corn;
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                                                                                                                                                                                                                                                                                                                   Claim 38; Page 85; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1997-202224/18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      n BE, rulland, AO, Skokut TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DOWELANCO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         O, Guo L,
Young SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2428
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pred. No.
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Zwick MG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hairpin ribozyme;
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Sequence

17

BP; 5

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ч; 0

RESULT 260 AAX01105/c ID AAX011 XX

AAX01105

standard; DNA; 17

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RASSULT 259
AAX84687/C
ID AAX846
XX AAX846
XX CAR AAX846
XX CONT 20-SEP
XX KW KDR SI
KW KDR SI
KW VASCULL
KW FEINO S
SYNTHE
OS THOMOS

XX TOOM
PA (SHIB)
XX WPI; 1
XX KDR SI
XX KDR SI
XX CC INVENT
OC CELL
CC INVENT
CC CINYENT
CC AND TO
CC INVENT
CC AND TO
CC ON TOOM
CC 
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Best Local
                                                                                                                Matches
                                                                                                                                    Best
                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KDR signal transduction inducer protein; human; diabetic receives vascular endothelial cell growth receptor; abnormal neovascularisation; kinase insert domain-containing receptor; solid tumour proliferation; gene therapy; metastasis; chronic rheumatoid arthritis; psoriasis; gene therapy; metastasis; chronic rheumatoid arthritis; psoriasis;
                                                                                                                                                                                                                                                                                                                                This sequence represents a PCR primer for DNA encoding the protein of the invention, which induces signal transduction of the vascular endothelial cell growth receptor KDR (kinase insert domain-containing receptor) by binding to its intracellular domain. The protein can be used in the investigation, diagnosis and treatment (including gene therapy) of diseases in which abnormal neovascularisation takes place, such as solid tumour proliferation, metastasis, chronic rheumatoid arthritis, psoriasis and retinopathy (including diabetic retinopathy and retinopathy of
                                                                                                                                                                                                                                                                         and retinopathy (including diabetic retinopathy and retinopathy prematurity). It may be used as a screen for candidate KDR signaturansduction inhibitors for therapeutic use.
                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX84687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 4; Page 67; 82pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KDR signal transduction inducing protein and antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shibuya M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primer for KDR signal transduction inducer protein coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-SEP-1999
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Local Similarity 76.9%;
hes 10; Conservative
                                                                                                                                       Pocar
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                                                      2262 GCGGCTTCCAGTC 2274
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15
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                                                                                                          l Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KYOWA HAKKO KOGYO KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; DNA;
                                                                                                                                                                                                                           17
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                                                                                                                                                                                                                        BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97JP-0343474.
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                                                                                                                                                                                                                        A,
                                                                                                                                                                                                                        5 C; 6
                                                                                                                                       100.0%;
                                                                                                                                                                    0.9%;
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                                                                                                          Score 13; DB; Pred. No. 2.3
0; Mismatches
                                                                                                             0;
                                                                                                                                                                                                                        G.
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Pred. No.
                                                                                                                                                                                                                           3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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0;
                                                                                                             2.3e+02;
hes 0;
                                                                                                                                                                    DB 1;
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                                                                                                                                                              Length 17;
                                                                                                                Indels
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                                                                                                                Gaps
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RESULT 261
ABN86972/c
ID ABN869
XX ABN869
XX ABN869
XX 29-JUL
XX 29-JUL
XX Prodru
KW Prodru
KW Simiar
KW Cancer
KW Cancer
KW famili
XX Cancer
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                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local !
                                                                                                                                                                                                                                                                                                                                         Matches
                            Simian virus 40; NS5B; viral infection; antiviral; cytostatic; HBV; antiallergic; immunosuppressive; gene therapy; AIDS; hepatitis B virus; cancer; leukaemia; genetic defect; allergy; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                agonists and antagonists; (ii) to raise antibodies; and (iii) therapeutically (reduced levels of AX are associated with resistance to nucleoside analogues with cytotoxic, anticancer and antiviral properties, with liver tumours, gout and acquired immune deficiency syndrome). Fragments of the DNA sequence are used as primers and probes to screen DNA libraries and for identifying AX-encoding nucleic acid, also as antisense therapeutics (particularly to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid encoding adenosine kinases and related oligo-nucleotides - expression vectors and transformed cells, used to modulate adenosine levels and to screen for specific modulators
                                                                      Prodrug ribozyme; ribozyme; SV40; HCV; hepatitis C virus; target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence is a PCR primer for DNA encoding the rat brain adenosine kinase (AK) of the invention. Cells transformed with the DNA are used produce recombinant AK. The AK is used: (i) to screen for specific
                                                                                                        Hepatitis C
                                                                                                                                          29-JUL-2002
                                                                                                                                                                        ABN86972;
                                                                                                                                                                                                    ABN86972 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cowart MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-JUN-1995;
                 familial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Column 43; 39pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-130392/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US5861294-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     adenosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Adenosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ocal adenosine concentrations at the site of tissue injury, increasing
                                                                                                                                                                                                                                                                                                            2400 GGAGGAACTTTTTAA 2414
                                                                                                                                                                                                                                                                                                                                             10;
                                                                                                                                                                                                                                                                                 16
                                                                                                                                                                                                                                                                                                                                                           Similarity
                genetic
                                                                                                                                                                                                                                                                               |:||:||:||:|
GRAGRAAYTTYTTRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                kinase; cytotoxic nucleoside resistance; anticancer; antiviral; our; gout; acquired immune deficiency syndrome; tissue injury; concentration; cytoprotection; rat; PCR primer; ss.
                                                                                                          virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Halbert DN,
                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A;
                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                       cytoprotection).
                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95US-0479614.
                disease;
                                                                                                        NS5B+ RNA oligonucleotide SEQ
                                                                                                                                                                                                                                                                                                                                                        0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               adenosine kinase coding
                                                                                                                                                                                                                                                                                                                                                                                                      2 C; 2 G; 4 T; 5 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kerwin JF, McNally T;
              primary genetic
                                                                                                                                                                                                                                                                                                                                           5
                                                                                                                                                                                                                                                                                                                                                         Score 13; DB 1;
Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                           Mismatches
                disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence.
                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                      Length 17;
                                                                                                          ID NO:10
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CC RNN-cleaving activity. Also described are: (i) an expression vector CC comprising (I) and preferably further comprising a tissue-specific promoter; (ii) a ribozyme prodrug comprising an intervening sequence in the ribozyme sequence removable by splicing, and lacking RNA-cleaving CC activity; (iii) a drug composition comprising (I); and (iv) the in vivo production of mature ribozyme with RNA-cleaving activity by introducing CC [I] into a eukaryote. (I) has antiviral, cycostatic, antializeric and immunosuppressive activities, and can be used in ribozyme and gene charapy. The ribozyme prodrug is useful e.g. in gene therapy. CC particularly for treating viral infections such as AlDS and those due to hepatitis B virus (HBV) and hepatitis C virus (HCV), cancers including those of the liver, pancreas and colon, and useamia, and diseases CC caused by genetic defects such as allergy, autoimmune diseases, familial genetic diseases and primary genetic diseases. The ribozyme prodrug, without RNA-cleaving activity, is encoded by a gene prodrug, converted the ribozyme sequence which can be spliced off in cytoplasm converted the ribozyme sequence which can be spliced off in cytoplasm converted the present invention.
Sequence 17 BP; 6 A; 5 C; 2 G; 4 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a gene (I) encoding a ribozyme prodrug comprising an intervening sequence removable by splicing, and/or lacking.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 81; 116pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel ribozyme prodrug without RNA-cleaving activity, for use e.g. i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     defective genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-256997/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SUMU) SUMITOMO PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yamamoto H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98JP-0249900.
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                                                                                                                                                                                                                                                                                                                                                                                                                  lacking
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S Matches Best Local Similarity Query Match 2469 GTACATGATGATG 2481 Conservative 0.9%; Score 13; DB 1; L; Pred. No. 2.3e+02; 0; Mismatches 0; 0, Length 17; Indels 0, Gaps 0

RRSULT 262
AA495774
ID AA4957
AC AA4957
XX
AC AA4957
XX
DF Human
XX
DE Human;
KW Human;
KW RNA cl
XX
DS Homo B
XX
PN WO2001
XX
PD 09-AUG
XX 밁 AAH95774 standard; RNA; 17 15 GTACATGATGATG 3 ₽P

Chk1 ribozyme substrate SEQ ID NO: 1199.

09-OCT-2001

(first entry)

Human; checkpoint kinase-1; Chk1; antisense; ribozyme; gene therapy; cleavage; cancer;

Homo sapiens

09-AUG-2001

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RESULT 263
ABK01806/c
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention provides nucleic acid molecules capable of downregulating the expression of the human checkpoint kinase-1 (Chk1) gene. These may be antisense or ribozyme sequences, and are useful in the treatment of diseases associated with conditions affected by Chk1 levels, including cancer. The present sequence is an oligonucleotide described in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene,
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                                                                                                                                                                                                                                                                                                                                          Homo sapiens
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17
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                                                   11-FEB-2000; 2000US-181797P.
28-FEB-2000; 2000US-185516P.
06-MAR-2000; 2000US-187128P.
                                                                                                                                                                     09-FEB-2001; 2001WO-US04273
                                                                                                                                                                                                                              16-AUG-2001
                                                                                                                                                                                                                                                                                       WO200159103-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                    Creutzfeldt-Jakob
                                                                                                                                                                                                                                                                                                                                                                                                                                                         chemotherapy-induced neuropathy; amyotrophic lateral sclerosis;
Parkinson's disease; ataxia; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human NOGO Zinzyme #128.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-MAR-2002
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(RIBO-) RIBOZYME PHARM INC
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Pred. No. 2.
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29-MAY-2002

(first entry) 17-mer scanning

ABN06895 standard; DNA;

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Human GDMLP-1

skeletal

Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine;

SEQ ID

NO:5

sequence

SEQ

heart disease; hGDMLP-1; heart; ID NO:6887.

myosin; chromosome 22; gene muscle disorder; amplicon;

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14

TCCAGCTCCTCCA 2

Matches Query Match Best Local

Similarity

100.0%;

Score 13; DB; Pred. No. 2.3

DB 1; ___ o. 2.3e+02; 0;

Length 17; Indels

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Gaps

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Conservative

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1573 TCCAGCTCCTCCA 1585

Sequence

17

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down comparison of a neurite growth inhibitor gene (NOGO).

The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CD NNAyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA molecule comparison of CD20 acids (e.g. a ribozyme or a colorif) pran amberzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with a NGN triplet), a zinzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with a NGN triplet), a zinzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving NA) with an NGN triplet), a zinzyme (cleaving NA) with an NGN triplet), a zinzyme (cleaving NA) with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the cuse of one or more therapies. In particular, the CD20 targetting cleaving nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma (NHL), bulky (cleaving NA) and inflammatory arthropathy. The NOGO-targetting cleaving nucleic acid is used to cleave RNA of the NOGO activity of the nucleic acid creat a patient having a condition associated with the level of NOGO. The creat apatient may further comprise the use of one or more therapies. Cleaving nucleic acid may be used to treat cation that is preferably myotrophic lateral sclerosis (MS), and/or chemotherapy-induced neuropathy, amotherapy and cerebrovascular accident (CVA, central nervous system (CNS) injury and cerebrovascular accident (CVA, charaese, muscular dystrophy, amotheraphic lateral sclerosis (MS), chambers which respond to the modulation of NOGO expression. The custes shich respond to the modulation of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BLAT/) BLATT L.
(MCSW/) MCSWIGGEN
(CHOW/) CHOWRIRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
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Homo sapiens

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                                                                                                                                                                                       CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 can be used as probes to detect, characterise CC and quantify hGDMLP-1 nucleic acids in samples, as amplification CC substrates, to provide initial substrates for the recombinant engineering CC of hGDMLP-1 protein variants having desired phenotypic improvements, and CC or expressing the proteins. The hGDMLP-1 proteins or polypeptides may CC be used as immunogens to raise antibodies that specifically recognise CC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption CC ionisation, as therapeutic supplement in patients having specific CC deficiency in hGDMLP-1 production, and in vaccines or for replacement CC diagnosing a disorder associated with the expression of hGDMLP-1, in CC particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the greening of the hGDMLP-1 sequence in the exemplification of the present
                                                      Query Match
Best Local S
Matches 13
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30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
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llarity 100.0%;
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2000US-234687P.
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format directly from WIPO
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RESULT 265

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CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of CC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 nucleic acids can be used as probes to detect, characterise CC and quantify hGDMLP-1 nucleic acids in samples, as amplification CC substrates, to provide initial substrates for the recombinant engineering CC of hGDMLP-1 protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may CC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP, proteins, as specific concentration and/or amount specifically of hGDMLP, proteins, as specific constation, as therapeutic supplement in patients having specific CC deficiency in hGDMLP-1 proteide sequences encoding hGDMLP-1 may be used for CC diagnosing a disorder associated with the expression of hGDMLP-1, in CC particular heart and skeletal muscle disorders. hGDMLP-1, in the cycening of the hGDMLP-1 sequence in the exemplification of the present
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invention.

N.B. The sequence data for this patent did not specification, but was obtained in electronic at ftp.wipo.int/pub/published_pct_sequence.
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21-SEP-2000;
27-SEP-2000;
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30-JAN-2001;
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2001WO-US00670.
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                                                                                                                                                             New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser description ionization, comprises human myosin-like protein hGDMLP-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the
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myosin-like

protein

proteins, or as specific biomolecule capture probes for surface enhanced laser desorption ionization, comprises

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New polypeptide, for raising antibodies that recognize hGDMLP-1

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RESULT 268
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Conservative (
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Query Match
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05-FEB-2001; 2001US-266860P.
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similarity 100.0%;
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RESULT 269
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; hear muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                 25-MAY-2001; 2001WO-US16981.
                                                                                                                                                                                                                                                                           WO200192524-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:9031.
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                                                                                                                                                                                                                                                                                                                                                         CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of GC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 nucleic acids can be used as probes to detect, characterise CC and quantify hGDMLP-1 nucleic acids in samples, as amplification of hGDMLP-1 protein variants having desired phenotypic improvements, and CC of hGDMLP-1 proteins. The hGDMLP-1 proteins or polypeptides may CC of hGDMLP-1 proteins. The hGDMLP-1 proteins or polypeptides may CC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption CC ionisation, as therapeutic supplement in patients having specific CC deficiency in hGDMLP-1 production, and in vaccines or for replacement CC therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for CC diagnosing a disorder associated with the expression of hGDMLP-1; in CC chromosome 22. The present sequence represents an oligomer used in the greening of the hGDMLP-1 sequence in the exemplification of the present convention.
                                                                                                                                                                                                                      Matches
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21-SEP-2000;
27-SEP-2000;
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                                                                                                                                                                                                                                                                                                         specification, at ftp.wino in
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Human GDMLP-1 17-mer scanning

SEQ ID NO:5 sequence SEQ ID NO:9032

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2397 CGTGGAGGAACTT 2409

Query Match Best Local S Matches 13

l Similarity

Conservative

100.0%; 0.9%;

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of CC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 uncleic acids can be used as probes to detect, characterise CC and quantify hGDMLP-1 nucleic acids in samples, as amplification CC substrates, to provide initial substrates for the recombinant engineering CC of hGDMLP-1 protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may CC be used as immunogens to raise antibodies that specifically recognise CC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific CC deficiency in hGDMLP-1 proteins for surface-enhanced laser description CC diagnosing a disorder supplement in patients having specific CC diagnosing a disorder associated with the expression of hGDMLP-1, in CC particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the convention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1 -
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30-JAN-2001;
30-JAN-2001;
                                                                             invention.

N.B. The sequence data for this patent did not specification, but was obtained in electronic f
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05-FEB-2001;
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cc hGDMLP-1 can be used in gene therapy and vaccine production. The characteristic characteristic characteristic characteristic cand quantify hGDMLP-1 nucleic acids in samples, as amplification consists to provide initial substrates for the recombinant engineering cc substrates, to provide initial substrates for the recombinant engineering cc of hGDMLP-1 protein variants having desired phenotypic improvements, and cf or expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise concentration and/or amount specifically of hGDMLP proteins, as specific concentration and/or amount specifically of hGDMLP proteins, as specific concentration and/or amount specifically of hGDMLP proteins, as specific constation, as therapeutic supplement in patients having specific confisation, as therapeutic supplement in patients having specific confisation.

CC deficiency in hGDMLP-1 production, and in vaccines or for replacement in patients having specific confisation, as therapeutic supplement in patients having specific confisation.
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                    The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGMMLP-1 nucleic acids can be used as probes to detect, characterists.
                                                                                                                                 Disclosure;
                                                                                                                                                                                              New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human
                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001;
30-JAN-2001;
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27-SEP-2000;
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Pred. No.
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and quantify hGDMLP-1 nucleic acids in samples, as amplification

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ARESULT 273
AREGORA 273
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ID AREGORA
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                                                                                                                                                                                                                                                                                    Diagnosing and/or determining a predisposition to a cellular proliferative disorder of breast tissue, in particular breast by determining the state of methylation of one or more nuclei
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    The sequence data for this patent did not form part of the printed
scification, but was obtained in electronic format directly from WIPO
ftp.wipo.int/pub/published_pct_sequence.

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llarity 100.0%;
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                                                                                                                                                                                                                                                                                        nucleic acids
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                                                 with a state of methylation from a subject not having the cellular proliferative disorder of breast tissue is indicative of a cellular proliferative disorder of breast tissue in the subject. The nucleic acids may be TWIST, HOXAS, NES-1, retinoic acid receptor beta (RARbeta), cestrogen receptor, cyclin D2, Wilms' tumour gene (WT-1), 14.3.3 sigma, HIN-1 or RASSFIA. The method is useful for diagnosing and/or determining a predisposition to a cellular proliferative disorder, in particular breast cancer including ductal carcinoma in situ, lobular carcinoma, colloid carcinoma, tubular carcinoma, medulary carcinoma, metaplastic carcinoma, intraductal carcinoma in situ, lobular carcinoma in situ and papillary carcinoma in situ. The present sequence is a primer used in the exemplification of the invention.
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Sequence 16 B₽; 2 A; 0 C; 8 G; 6 T; 0 other;

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S
               Matches
                       Query Match
Best Local
              14;
                       Similarity
 CAGCCAACTGCACCAA 2462
              Conservative
                      0.9%;
              0,
                      Score 12.8;
Pred. No. 2
                Mismatches
                     .3e+02;
                               В
                             ۳.
                             Length
                Indels
                0
               Gaps
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0

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16 CACCCAACTTCACCAA 1

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RESULT 274
ABA89557
ID ABA895
XX ABA895
XX ABA895
XX ACido
XX Seria
XX Seria
XX Geoch
XX Geoch
XX Geoch
XX Ho-2
PN WO20
XX ACido

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABA89557;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                environmental analysis; parasite detection; virus detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Acidobacterium Tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
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18-OCT-2001.

10-APR-2001; 2001WO-US11609

10-APR-2000; 11-APR-2000; 2000US-196258P. 2000US-196063P.

(ASHB/) ASHBY M.

WPI; 2002-010926/01.

Determining genetic diversity of population by analyzing a specific polymorphic region characteristic of particular genome in population interest, useful for locating mineral deposits or petroleum reserves , ef

Disclosure; Page 50; 83pp; English.

The present invention relates to a method of determining the genetic diversity of a population, involving amplifying a genome subregion with a polymorphic site, cleaving amplified fragment close to the polymorphic site, immobilising the amplified fragment, splitting into two pools, adding a linker to each pool, digesting the immobilised product to form tags that are ligated to form diags, and amplifying, cleaving and ligating to form concatemers and sequencing. The method is known as serial analysis of ribosomal DNA (SARD). This can be used to determine the genetic diversity of a population including microbial, viral or immune cell populations. The microbial population whose genetic diversity can be determined is from a sample associated with a site for not contains. or natural gas exploration, i.e.,

proliferative disorder of by state of methylation of one subject, where the state of

methylation

of

compared

present invention relates to a metho

es to a method of diagnosing a cellular reast tissue, which involves determining the or more nucleic acids isolated from the

method of diagnosing issue, which involves

Claim 13;

Page

62; 115pp; English.

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RESULT 275
AAI68644
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Best Local :
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antiasthmatic, antiinflammatory, immunosuppressive and gastrointestinal activity. (A) are used for treatment or prevention of ICAM-1-associated diseases, specifically psooriasis, neurodermatitis, allergic asthma, Crohn's disease, autoimmune diseases and transplant rejection. Compared with antisense oligonuclectides. (A) provide a longer-lasting effect (they bind directly to the gene, so a compensatory increase in transcription is not possible). (A) may be coupled to psoralen to provide light-regulatable, sequence-specific downregulation of genes; this should
                                                                                                                                This invention describes novel polydeoxyribonucleotides (A), for use as triple-helix forming oligonucleotides, having at least 3 sequential purine and/or pyrimidine bases, capable of inhibiting transcription of ICAM-1. (A) has a sequence specific for the transcribed or promoter regions of the ICAM-1 (intracellular adhesion molecule) gene. The products of the invention have antipsoriatic, dermatological,
                                                                                                                                                                                                                                                                                          Triple-helix forming polydeoxyribonucleotides, useful for treating intracellular adhesion molecule-1 related diseases, e.g. psoriasis, directed against transcribed or promoter regions of the ICAM-1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   reserves, associated with a site of mineral exploration, associated with a agricultural field, of patient sample suspected to have bacterial or fungal infection, associated with bioremediation site, or of an insect or parasite. The methods have application in fields of geochemical exploration, agriculture, bioremediation, environmental analysis, clinical microbiology, forensic science and medicine. The present sequence is an oligonucleotide described in the exemplification of the
                                                                                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-017614/02
                                                                                                                                                                                                                                                                                                                                                                                                 Degitz KK,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      intracellular adhesion molecule; dermatological; antiasthmatic; antiinflammatory; immunosuppressive; gastrointestinal; psoriasis; neurodermatitis; allergic asthma; Crohn's disease; autoimmune disease; transplant rejection; psoralen; photo-ultra-violet therapy; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ICAM-1 triple helix associated oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                    (BESC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             [CAM-1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                DEGITZ
BESCH F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           triple helix; transcription inhibition; antipsoriatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                            Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CGGCTGTGCCGAAGCT 16
                                                                                                                                                                                                                                                                                                                                                                                                  Beach
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                                                                                                                                                                                                                                                              14;
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                                                                                                                                                                                                                                                         61pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C; 6 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16
                                                                                                                                                                                                                                                              German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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Pred. No. 2.3e+
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2;
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RESULT 276
AA168645/C
ID AA1686
This invention describes novel polydeoxyribonucleotides (A), for use as C triple-helix forming oligonucleotides, having at least 3 sequential CC purine and/or pyrimidine bases, capable of inhibiting transcription of CC ICAM-1. (A) has a sequence specific for the transcribed or promoter of the ICAM-1 (intracellular adhesion molecule) gene. The CC products of the invention have antipsoriatic, dermatological, CC antiasthmatic, antinflammatory, immunosuppressive and gastrointestinal CC activity. (A) are used for treatment or prevention of ICAM-1-associated CC diseases, specifically psoriasis, neurodermatits, allergic asthma, CC Crohn's disease, autoimmune diseases and transplant rejection. Compared CC with antisense oligonucleotides, (A) provide a longer-lasting effect (they bind directly to the gene, so a compensatory increase in CC transcription is not possible). (A) are observable to psoralen to provide light-regulatable, sequence-specific downregulation of genes; this should CC make photo-ultra-violet therapy more specific, with reduced side effects. AxI68599-AXI6873 represent oligonucleotides used to illustrate the
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Best Local
                                                                                                                                                                                                                                                                                                                                                                Triple-helix forming polydeoxyribonucleotides, useful for treating intracellular adhesion molecule-1 related diseases, e.g. psoriasis, directed against transcribed or promoter regions of the ICAM-1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             make photo-ultra-violet therapy more specific, with reduced side effects. AAI68599-AAI68673 represent oligonucleotides used to illustrate the
                                                                                                                                                                                                                                                                                                                                  Claim 5; Page 15; 61pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-017614/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Degitz KK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200179487-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       intracellular adhesion molecule; dermatological; antiasthmatic; antiinflammatory; immunosuppressive; gastrointestinal; psoriasis; neurodermatitis; allergic asthma; Crohn's disease; autoimmune disease; transplant rejection; psoralen; photo-ultra-violet therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-JAN-2002
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C;
                                                                                                                                                                                                                                                                                                                                    German
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Pred. No. 2.
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Sequence

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RESULT 277
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Best Local Similarity
                                                     Matches
                                                                               Query Match
                                                                                                                                The invention relates to a novel isolated nucleic acid molecule comprising a variant gene associated with an eating disorder and selected from any of 119 polymorphisms with their corresponding genotyping in dataset, alleles and HGBASE identification, given in the specification. The novel nucleic acid molecule has polymorphisms in the serotonin receptor 1D, delta-ophoid receptor, or dopamine receptor D2, which is useful in diagnostic and prognostic assays for eating disorders, in particular anorexia nervosa and bulimia nervosa. This polymucleotide sequence represents a dopamine D2 receptor PCR primer of the invention.
                                                                                                                                                                                                                                                                                                      New nucleic acid molecule having polymorphisms in the serotonin receptor 1D, delta-optoid receptor, or dopamine receptor D2, useful in diagnostic and prognostic assays for eating disorders, such as anorexia and bulimia nervosa
                                                                                                          Sequence 16 BP; 0
                                                                                                                                                                                                                                                                         Example 2; Page 41; 149pp; English.
                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-268122/26.
                                                                                                                                                                                                                                                                                                                                                                                                     Bergen AW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-NOV-2001;
19-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-JUL-2001;
20-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-JUL-2002; 2002WO-US22555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003012143-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      anorexia nervosa; bulimia nervosa; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eating disorder; polymorphism; dataset; allele; HGBASE identification; serotonin receptor 1D; delta-opioid receptor; dopamine receptor D2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dopamine-D2-receptor PCR primer SEQ ID No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABT34223;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABT34223 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                 (PRIC-) PRICE FOUND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                            L9-DEC-2001;
                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2421 GAAGGAAGGACACAGA 2436
                     1785 CAAAGACAAGCCCAAG 1800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16 GAAGGAAGGAAAAAAA 1
16
                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; 2001US-305153P.
; 2001US-306440P.
; 2001US-331285P.
; 2001US-340843P.
                                                  0.9%;
ilarity 87.5%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     Yeager M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001US-340844P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                         A; 2 C; 7 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.9%;
87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12.8; DI
Pred. No. 2.3e
0; Mismatches
                                                   0
                                                                Score 12.8; DB Pred. No. 2.3e+(
                                                  Mismatches
                                                                2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9
                                                                            Length 16;
                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                   0
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RESULT 278

RNA; enzyme; enzymatic RNA molecule; ERM; cleave; RNA; mRNA; h picornavirus; HIV; immunodeficiency virus; hepatitis B virus; papilloma virus; HPV; Epstein-Barr virus; EBV; TCLV; T-cell leukaemia virus; hepatitis C virus; HCV; cytomegaloviru influenza virus; HSV; herpes simplex virus; vector; immune res

HBV;

s; hepatitis C virus; HCV; cytomegalovirus; herpes simplex virus; vector; immune respo

treatment;

83

Herpes simplex virus target sequence

21.

antibody; ribozyme; viral RNA;

Synthetic.

WO9323569-A1

29-APR-1993; 25-NOV-1993.

93WO-US04020

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                                                                                                                                                                                   RESULT 279
AAQ52943
                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                            Probe A1
(Updated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAN91716;
                                                                                                                                                                                                                                                                              Sequence 17
                                                                                                                                                                                                                                                                                                                   Disclosure;
                                                                                                                                                                                                                                                                                                                                 Toxin protein of Bacillus thuringiensis bacteria - using a potentiating amt. of a trypsin inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    EP339009-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
14-MAR-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAN91716 standard;
                                                                                                                                      25-MAR-2003
26-MAY-1994
                                                                                                                                                             AAQ52943;
                                                                                                                                                                           AAQ52943 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                         WPI; 1989-311431/43.
                                                                                                                                                                                                                                                                                                                                                                      Fuchs RL, Kishore GM,
                                                                                                                                                                                                                                                                                                                                                                                                      11-APR-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                     14-APR-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                    25-OCT-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bacillus thuringiensis tenebrionis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Probe A1; Bacillus thuringiensis tenebrionis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Probe Al for Bacillus thuringiensis tenebrionis.
                                                                                                                                                                                                                                                                                                                                                                                       (MONS ) MONSANTO
                                                                                                                                                                                                                                 1585
                                                                                                                                                                                                                                                14;
                                                                                                                                                                                                                   \vdash
                                                                                                                                                                                                                                ATGAACTCCAACACCC 1600
                                                                                                                                                                                                                                                                                            on
si
                                                                                                                                                                                                                   ATGAACCCCAACAACC 16
                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                              BP; 7 A; 7 C; 2 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                            based on amino acids 1-6 of the B.t.t. 25-MAR-2003 to correct PF field.
                                                                                                                                                                                                                                                                                                                  page 13; 56pp; English.
                                                                                                                                      (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (updated)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                     89EP-0870047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                      0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17
                                                                                                                                                                                                                                                                                                                                                                         MacIntosh
                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                °.
                                                                                                                                                                                                                                                       Score 12.8; DB 1;
Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                         sc;
                                                                                                                                                                                                                                                              Length 17
                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                          improved in efficacy
                                                                                                                                                                                                                                                                                                     toxin
                                                                                                                                                                                                                                                                                                     gene
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                                                                                                                                                                                                                                                Gaps
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RESULT 280
AAX64062
ID AAX640
XX
AC AAX640
XX
DT 20-JUL
XX
DE Rabbit
XX
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                                                                                                                                                                     Matches
                                                                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                                                             complementary to a substrate binding region in the specified gene target. They also have enzymatic activity, in that they specifically cleave RNA in the target. The ERMs interfere with viral replication a therefore have anti-viral properties. They can be used to attenuate viruses to be used in vaccines. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-MAY-1992;
14-MAY-1992;
14-MAY-1992;
          Rabbit stromelysin hammerhead target SEQ ID NO:694
                                                   AAX64062
                                                                                                                                                                                                            Sequence 17
                                                                                                                                                                                                                                                                                                                 The sequences (AAQ52923-Q53037) are pref. herpes simplex virus target sequences for enzymatic RNA molecules. The RNA molecules are
                               20-JUL-1999
                                                                       AAX64062 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                     infection
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                                                                                                                                                                                                                                                                                                                                                                                                                        Draper KG,
Mamone JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-AUG-1992;
26-AUG-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-MAY-1992
31-JUL-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-MAY-1992
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14-MAY-1992
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14-MAY-1992
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14-MAY-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RIBO-) RIBOZYME PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-SEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-MAY-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          L5-OCT
                                                                                                                                              2279 GGATGGCTCCAGAAGC 2294
                                                                                                                                                                   al Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                5; Fig
                                                                                                                           N
                                                                                                                           GGGUGGCUCCAGAACC 17
                                                                                                                                                                                                                                                                                                                                                                   RNA molecules - used to and gene expression
                                                                                                                                                                                                            BP; 3
                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                 Dudycz LW,
                                                                                                                                                                                                                                                                                                                                                 15;
                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92US-088288

92US-088289

92US-0882921

92US-0882922

92US-0883823

92US-0884073

92US-0884074

92US-0884431

92US-0884431

92US-0884431

92US-0884431

92US-0884431

92US-0884431

92US-0884431

92US-0884431

92US-0884431

92US-088433

92US-088433

92US-088433

92US-088433

92US-088433
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92US-0882823.
92US-0882824.
92US-0882886.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          92US-0987133
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                                                                                                                                                                                                                                                                                                                                                287pp; English
                                                                                                                                                                                                            A; 5
                                                                                                                                                                           0.9%;
                                                                                                                                                                                                            C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                       INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Mcswiggen JA,
                                                                                                                                                                                                            G; 2
                                                                                                                                                                   2;
                                                                                                                                                                            Score
Pred.
                                                                                                                                                                   Mismatches
                                                                                                                                                                                                            U; 0 other;
                                                                                                                                                                              No. 2.5e+0
                                                                                                                                                                                                                                                                                                                                                                               inhibit viral
                                                                                                                                                                  8; DB 1;
2.5e+02;
ches 2;
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                                                                                                                                                                                                                                                                                                                                                                              replication
                                                                                                                                                                                     Length 17;
                                                                                                                                                                    Indels
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                                                                                                                                                                   0
                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                    and
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RESULT 281 AAX64063 ID AAX640

AAX64063 standard; RNA; 17

믕 S

N

17

1694

GGGAGTTTCCAAGAGA 1709

Query Match Best Local S Matches 12

l Similarity 12; Conserv

Conservative

2;

Mismatches

Indels

0,

Gaps

0

0.9%;

Score 12.8; Pred. No. 2

2.5e+02;

DB

1; <u>ب</u>

Length 17;

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The present invention describes a novel enzymatic nucleic acid (ENA)
Chaving a hammerhead motif (EM) comprising: (i) at least 5 ribose
residues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii)
at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification.
Che ENA; can inhibit collagenase and stromelysin production in the
synovial membrane of joints for the treatment or prevention of arthritis,
particularly osteoarthritis or rheumatoid arthritis. The ENA's can also
be used to treat antigen presenting cells of a donor to induce tolerance
in a recipient to an alloantigen of a donor. They can also be used for
chancing graft tolerance or for treating autoimmune disease, and for
treating allergies and other inflammatory conditions. The ENA's can also
be used in diagnosis. Ribozyme therapy impacts on the expression of
stromelysin without introducing the non-specific effects upon gene
expression which accompany treatment with retinoids and dexamethasone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Draper K,
Beigelman
Sequence
                                             The concentration of ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly specific. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-MAY-1995;
04-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oryctolagus cuniculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Arthritic condition; graft tolerance; immune response; target; cleavage; hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatiod arthritis; autoimmune disease; allergy; inflammation;
                                 present invention.
                                                                                                                                                                                                                                                                                                                                         Example 1; Page 156; 307pp; English
                                                                                                                                                                                                                                                                                                                                                                                        Enzymatic nucleic acid molecules having a for the treatment of arthritis, induction
                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1996-300653/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matulic-Adamic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-APR-1995
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23-DEC-1994;
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13-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-NOV-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RIBOZYME PHARM INC.
 17
                                                                                                                                                                                                                                                                                                                                                                          of auto-immune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ŗ
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L, Karpeisky A, N
amic J, Jarvis T,
 BP,
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94US-
94US-
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95US-0000974.
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95US-0426124.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-0512861.
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95US-0434509.
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Α;
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-0354920.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -0363253.
4 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   McSwiggen J, Pavcc
A, Modak A, Usman
s T, Thompson JD,
6 G;
2 U;
0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pavco P, Stinchcon Usman N, Burgin A; JD, Wincott F;
                                                                                                                                                                                                                                                                                                                                                                                      hammer-head motif - us
of graft tolerance or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Stinchcomb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ĎŢ,
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Ś
                              Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                               The present invention describes a novel enzymatic nucleic acid (ENA) having a hammerhead motif (HM) comprising: (i) at least 5 ribose residues; (ii) a 2'-C-allyl modification at position 4 of the ENA, (iii) at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification. The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of adonor. They can also be used for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-APR-1995;
02-MAY-1995;
04-MAY-1995;
07-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-DEC-1994;
23-DEC-1994;
17-FEB-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Arthritic condition; graft tolerance; immune response; target; cleavage; hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; collagenase stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                        Sequence 17
                                                                                                                                   The concentration of ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly specific. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 156; 307pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                             treatment of auto-immune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Enzymatic nucleic acid molecules having a hammer-head motif - unifor the treatment of arthritis, induction of graft tolerance or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1996-300653/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matulic-Adamic J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Beigelman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-NOV-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9618736-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rabbit stromelysin hammerhead target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-JUL-1999
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1694 GGGAGTTTCCAAGAGA 1709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RIBOZYME PHARM INC
                                                                                                                         invention.
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L, Karpeisky A,
                            0.9%;
nilarity 75.0%;
Conservative
                                                                                          BP; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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95US-0432874.
95US-0434509.
95US-0000951.
95US-0000974.
95US-0512861.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jarvis T,
                                                                                          σı
                                                                                          C; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 McSwiggen J, Pavco P, Stinch
A, Modak A, Usman N, Burgin
a T. Thompson JD, Wincott F;
                                                                                          G; 2
                              2
                            Score 12.8; DB
Pred. No. 2.5e+(
2; Mismatches
                                                                                          U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO:695
                                            2.5e+02;
                                                          1:
                                                        Length 17;
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RESULT 283
AAT81588/c
ID AAT815
XX
AC AAT815
XX
DT 21-DEC
XX
DE Human
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AAT81587/c
                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                     The present sequence represents the preferred target sequence for an enzymatic nucleic acid, especially a hammerhead ribozyme, which cleaves the human c-myb sequence at the base position indicated in the descriptor line. The c-myb sequence was screened for optimal ribozyme target sites using a computer folding algorithm, and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes cleave the c-myb sequence and can be used to prevent smooth muscle cell hyperproliferation in restenosis, especially after coronary angioplasty, and in cancers.
Human c-myb hammerhead ribozyme target sequence (nt. position 2972).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New enzymatic nucleic acid molecules - which cleave RNA produced e.g. c-myb, for treating restenosis or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic nucleic acid; hammerhead; ribozyme; cleavage; human; smooth muscle cell; hyperproliferation; restenosis; cancer; c-myb; coronary angioplasty; ss.
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                              21-DEC-1997
                                                                                                                                                                                                                                                                               Sequence 17 BP; 1 A; 0 C; 4 G; 12 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 79; 128pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Draper K, Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human c-myb
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                                                                                      AAT81588
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18-MAY-1994;
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                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                          2240 ATTACAAAAAGACCAC 2255
                                                                                                                                                                                                                       14;
                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGGAGCUUCCACGAGA 16
                                                                                                                                                              ATAACAAAAAAACCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hammerhead ribozyme target sequence (nt. position 2971).
                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                            (first entry)
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94US-0245466.
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87.5%;
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                                                                                      RNA;
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Pred. No. 2.5e+02;
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AAT81253
ID AAT8125
XX AAT812
AC AAT812
XX O5-OCT
XX DF Human
XX Enzyma
KW Enzyma
KW Encymb
KW C-myb;
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Best Local S
Matches 14
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                                                                                                                                    Enzymatic nucleic acid; hammerhead; ribozyme; cleavage; human; smooth muscle cell; hyperproliferation; restenosis; cancer; c-myb; coronary angioplasty; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New enzymatic nucleic acid molecules -
                                                                                                                                                                                                Human c-myb hammerhead ribozyme target sequence (nt. position 1587)
                                                                                                                                                                                                                                05-OCT-1997
                                                                                                                                                                                                                                                                 AAT81253;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 79; 128pp; English.
                18-MAY-1995;
                                              23-NOV-1995.
                                                                            WO9531541-A2
                                                                                                                                                                                                                                                                                              AAT81253 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                coronary angioplasty, and in cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1996-010927/01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC.
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18-MAY-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     c-myb; coronary angioplasty; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Enzymatic nucleic acid; hammerhead; ribozyme; cleavage; human;
smooth muscle cell; hyperproliferation; restenosis; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                          2240
                                                                                                         sapiens
                                                                                                                                                                                                                                                                                                                                                                            16
                                                                                                                                                                                                                                                                                                                                                                                                                                         14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                         ATTACAAAAAGACCAC 2255
                                                                                                                                                                                                                                                                                                                                                                            ATAACAAAAAAACCAC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                       0.9%;
larity 87.5%;
Conservative
                                                                                                                                                                                                                                (first entry)
                95WO-US06368
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94US-0245466.
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Pred. No. 2.5e+02;
0; Mismatches 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 which cleave RNA produced
                                                                                                                                                                                                                                                                                                                                                                                                                                       .5e+02;
les 2;
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RESULT 285
AAT81188/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents the preferred target sequence for an enzymatic nucleic acid, especially a hammerhead ribozyme, which cleaves the human c-myb sequence at the base position indicated in the descriptor line. The c-myb sequence was screened for optimal ribozyme target sites using a computer folding algorithm, and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme esites were identified. Ribozymes were synthesised and their activities optimised by either varying the length of the binding
Claim 1; Page 68; 128pp; English
                        New enzymatic nucleic acid molecules - which cleave e.g. c-myb, for treating restenosis or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT81188 standard; RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               arms or by modification to prevent degradation by nucleases. The ribozymes cleave the c-myb sequence and can be used to prevent smooth muscle cell hyperproliferation in restenosis, especially after
                                                                    WPI; 1996-010927/01.
                                                                                            Draper K, Jarvis T,
                                                                                                                                                  13-JAN-1995;
18-MAY-1994;
                                                                                                                                                                                                                     23-NOV-1995.
                                                                                                                                                                                                                                                 WO9531541-A2
                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                     c-myb;
                                                                                                                                                                                                                                                                                                                                Enzymatic nucleic acid; hammerhead; ribozyme; cleavage; human;
                                                                                                                                                                                                                                                                                                                                                          Human c-myb hammerhead ribozyme target sequence (nt. position 1257).
                                                                                                                                                                                                                                                                                                                                                                                        29-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                   AAT81188;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   coronary angioplasty, and in cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New enzymatic nucleic acid molecules - which cleave RNA produced
                                                                                                                         (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                          18-MAY-1995;
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18-MAY-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13;
                                                                                                                                                                                                                                                                                                     coronary
                                                                                                                                                                                                                                                                                                                  muscle cell; hyperproliferation; restenosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for treating restenosis or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                 95US-0373124.
94US-0245466.
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94US-0245466.
                                                                                                                                                                                                                                                                                                     angioplasty; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%;
                                                                                           McSwiggen
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Pred. No. 2.5e+02;
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                                       RNA produced by
                                                                                                                                                                                                                                                                                                                    cancer;
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RRESULT 286
RAX75356/C
ID AAX753
XX AAX753
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XX Vascul
DT 28-JUL
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                                                        The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour
   angiogenesis,
be treated by
                                                                                                                                                                                                                                                                                                        Claim 4;
                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1997-259017/23
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(RIBO-)
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26-OCT-1995;
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tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease;
fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse flt-1 VEGF receptor hammerhead ribozyme substrate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 foetal liver
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treated by
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                               psoriasis and rheumatoid
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molecule or the expression
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                                                                            the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGP). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psocriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples
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Sequence
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26-OCT-1995;
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                                                        of nucleic acid molecules from the present invention.
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(RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-OCT-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; Page 138; 218pp; English.
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                                                                  Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1, KDR, hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular diseafms-like tyrosine kinase 1; kinase insert domain containing recept foetal liver kinase 1; ss.
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26-OCT-1995;
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                                                                                     domain containing receptor;
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WO9715662-A2 Homo sapiens.

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RESULT 291
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                          (CHIR )
(RIBO-)
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Similarity 87.5%;
14; Conservative
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                             RIBOZYME PHARM INC
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Pred. No. 2.5e+02;
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Mismatches 2;
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RRESULT 292
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KW Fins-1;
KW foetal
XX Lumour
KW fms-1;
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12; Conserv
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Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient

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WPI; 1997-259017/23.

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McSwiggen

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Pavco

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Stinchcomb

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(CHIR) (RIBO-)

RIBOZYME PHARM INC. CHIRON CORP

The present invention describes nucleic acid molecules which modulate

Claim 4;

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62; 218pp; English

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RESULT 293
AAT47828/c
ID AAT478
XX AAT478
XX AAT478
XX DT 14-MAY
XX NF2; n
KW bilate
KW mening
KW diagno
KW PCR; p
XX Synthe
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                   AAT47828-T47830 are PCR primers used for the isolation of the NF2 (neurofibromatosis type 2) gene from various murine tumours. NF2 is an autosomal, dominantly inherited disorder characterised by multiple tumours of the central nervous system, predominantly bilateral vestibular schwannomas (acoustic neuromas) of the eighth cranial nerve. Other symptoms of NF2 include cranial meningiomas, spinal nerve root schwannomas and presentle lens opacities. The NF2 gene, mapped to chromosomal region 22q12 between the loci D22S1 and D22S28, acts a tumour suppressor. The NF2 gene is alternatively spliced resulting in three different isoforms encoding three differing functions. Merlin isoforms I-III, which are likely to have differing functions. Merlin stands for moesin-ezrin-radixin like protein, so called due to enther articles and protein sections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NF2; neurofibromatosis type 2; multiple tumours; nervous system; bilateral vestibular schwannoma; acoustic neuroma; cranial nerve; meningioma; lens opacity; chromosome region 22q12; tumour suppressor; merlin; moesin-erzin-radixin like protein; alternative splicing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             develop
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1997-020406/02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bianchi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-JAN-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BRIM ) BRISTOL-MYERS SQUIBB CO
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prods. for
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Column 14; 46pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kley NA,
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the
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Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BR.
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neurofibromatosis
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Query Match Best Local Matches

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Matches
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                                                  The present invention describes enzymatic nucleic acid molecules (NAMs) which specifically cleave RNA derived from an epidermal growth factor receptor (EGF-R) gene. AAV97221 to AAV98043 and AAV98979 to AAV99090 represent specifically claimed target sequence from human EGF-R. AAV98044 to AAV98866 and AAV98867 to V9978 represent hammerhead ribozymes and hairpin ribozymes respectively for human EGF-R. The NAMs are useful for cleaving EGF-R RNA in the treatment of a condition associated with EGFR expression levels e.g. to inhibit cell proliferation in the prevention or treatment of cancers. The NAMs can also be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of EGF-R RNA in a cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-DEC-1997;
31-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                     Claim 5; Page 74; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Enzymatic nucleic acids - which cleave RNA derived from an epidermal growth factor receptor, useful for inhibiting cell proliferation and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-437449/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Akhtar S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer; genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; epidermal growth factor receptor; EGFR; EGF-R; target sequence; hammerhead ribozyme; hairpin ribozyme; inhibition; cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 6 A; 1 C; 7 G; 3 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and proteins encoded by them, are useful in diagnosing NF2 disea Merlin protein products act as tumour suppressors and can be use suppress tumour growth, as can the cDNA sequence in gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                              treating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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97US-0036476.
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AC AAV946
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24-FEB
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                                                                                                                                                                                                                         RESULT 295
AAV97404/c
ID AAV97404 standard; RNA; 17 BP
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                                                                                                                                                                     Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                           The present invention describes enzymatic nucleic acid molecules (NAMs) which specifically cleave RNA derived from an epidermal growth factor receptor (EGF-R) gene. AAV97221 to AAV98043 and AAV98979 to AAV99909 represent specifically claimed target sequence from human EGF-R. AAV98040 to AAV98066 and AAV98067 to V9878 represent hammerhead ribozymes and hairpin ribozymes respectively for human EGF-R. The NAMs are useful for cleaving EGF-R RNA in the treatment of a condition associated with EGFR expression levels e.g. to inhibit call proliferation in the prevention or treatment of cancers. The NAMs can also be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of EGF-R RNA in a cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; epidermal growth factor receptor; EGFR; EGF-R; target sequence; hammerhead ribozyme; hairpin ribozyme; inhibition; cell proliferation; cancer; genetic drift; detection; mutation; ss.
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 24-FEB-1999
                           AAV94632;
                                                   AAV94632 standard;
                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                          Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                   for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzymatic nucleic acids - which cleave RNA derived from an epidermal growth factor receptor, useful for inhibiting cell proliferation and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-437449/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Akhtar S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-DEC-1997;
31-JAN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYAS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-)
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                                                                                                                                           1358 CGCCTGGAAGAAAA 1373
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                                                                                                                                                                                 o.9%;
Similarity 87.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGTGATGTCTGGTCCT 2338
                                                                                                                                                                                                                         17
                                                                                                                    CGACTGCAAGAGAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                         Page 71; 109pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGUGAUGUCUGGAGCU 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fell P,
                                                                                                                                                                                                                       BP; 1
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                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                     cancers
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 (first entry)
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97US-0036476.
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                                                                                                                                                                                                                         A; 4 C; 3
                                                   RNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                         English.
                                                    ВÞ
                                                                                                                                                                                                                         G; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleotide position
                                                                                                                                                                    Pred. No. 2.5e
0; Mismatches
                                                                                                                                                                                 Score 12.8; DB 1;
Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JA;
                                                                                                                                                                                                                         u;
0
                                                                                                                                                                                                                         other;
                                                                                                                                                                      1.5e+02;
1es 2;
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                                                                                                                                                                                               Length 17;
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RESULT 297
AAV96482/c
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Best Local Similarity
Matches 14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 4 A; 4 C; 3 G; 6 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence invention describes ribozymes targeted to modulate the synthesis and/or expression of interleukin (II)-2R gamma encoded RNA. AAV93889 to AAV94574 represent specifically claimed ribozymes, and AAV94575 to AAV95260 represent specifically claimed substrate sequences from the present invention. The ribozymes can be used for the treatment of, e.g. graft rejection, autoimmune disease, cancer, psoriasis, allergy and other inflammatory conditions. The ribozymes are also used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ribozymes targeted to interleukin 2 - useful for treating e.g. cancer, autoimmune disease and allergies
24-NOV-1997;
28-JAN-1997;
                                                                                                                                                               Solanidine; glucosyltransferase; potato; citrate synthase; target; hammerhead ribozyme; hairpin ribozyme; alkaloid biosynthesis;
                                         14-JAN-1998;
                                                                  30-JUL-1998
                                                                                                                        Solanum tuberosum
                                                                                                                                                                                                      Potato citrate synthase target sequence position
                                                                                                                                                                                                                                  01-MAR-1999
                                                                                                                                                                                                                                                                                      AAV96482 standard; RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           to induce tolerance in a recipient to alloantigen from a donor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4;
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                                                                                             WO9832843-A2
                                                                                                                                                 flower formation; cleavage; solanaceous plant; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                              2156 CAGCCAGAAATGTTTT
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                                                                                                                                                                                                                                                                                                                                                         CAGCCAGAAGTGATTT 1
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                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                 (first entry)
97US-0979416.
97US-0036545.
                                         98WO-US00738.
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                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Pred. No. 2.5e+02;
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                                                                                                                                                                                                        546
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RESULT 298
AAA18720/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                       Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot wine stain; Sturge Weber syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            solanaceous plants, particularly potato but also tomato, pepper, aubergine and ditura or to inhibit flowering in potato, lettuce, spinach, cabbage, brussel sprouts, arugula, kale, collards, chard, beet, turnip, sweet potato and turf grass. Also the ribozymes can be used for RNA manipulation in the same way that restriction endonucleases are for DNA, as well as to examine genetic drift and mutations in plants and to detect specific RNA. The ribozymes can be targeted to specific genes or to consensus sequences within a family of related genes, and being catalytic need to be present at only very low concentrations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RNA-cleaving activity (e.g. ribozymes) which are capable of modulating the expression of plant genes: (i) involved in biosynthesis of alkaloids, or (ii) involved in flower formation. AAV9592 to AAV96334, and AAV96335 to AAV96354 represent potato solanidine glucosyltransferase hammerhead and hairpin ribozymes, respectively. AAV95629 to AAV95931, and AAV96335 to AAV96734 represent potato solanidine glucosyltransferase target sequences. AAV96737 to AAV97170, and AAV97171 to AAV97195 represent potato citrate synthase hammerhead and hairpin ribozymes, respectively. AAV96735 to AAV96772, and AAV97196 to AAV97220 represent potato citrate synthase target sequences.Ribozymes of the present potato citrate synthase target sequences.Ribozymes of the present invention can be used to inhibit the synthesis of toxic alkaloids in solanoceous nlants, natricularly potato.
24-MAR-1999;
                                       07-OCT-1999.
                                                                                                                  Homo sapiens
                                                                                                                                                       Kippel-Trenaunay-Weber
                                                                                                                                                                                                                                                                                                                                                                                                   19-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA18720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA18720 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     biosynthesis or regulating flowering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New enzymatic nucleic acid(s) - useful for,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-427939/36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2468 TGTACATGATGAG 2483
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TATACATGATGATCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 53; 79pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 5
                                                                                                                                                                                                                                                                                                                                                          substrate sequence SEQ ID NO:1946.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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99WO-US06507.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; 4 C; 2 G; 6 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.9%;
                                                                                                                                                         syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
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Pred. No. 2.5e+02;
                                                                                                                                                       Osler-Weber-Rendu syndrome; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e.g. reducing alkaloid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 17;
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Sequences for integrin alpha 6 subunit, and AAA23362 to AAA21500 and AAA21596 to AAA21688 represent their corresponding target sequences; AAA21698 to AAA22475 and AAA23263 to AAA2342 represent ribozyme sequence for integrin subunit beta 3, and AAA23476 to AAA23363 to AAA23432 represent their corresponding target sequences. The ribozymes of the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARRY, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2,
                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, and AAA17168 to AAA17560 and AAA17633 to AAA17684 represent their corresponding target sequences for Tie-2, AAA18385 and AAA18386 to AAA19087 to AAA19155 to AAA19222 represent their corresponding target sequences; AAA1923 to AAA19223 to AAA19222 represent their corresponding target sequences; AAA1923 to AAA1923 to AAA2351 and AAA19251 to AAA2351 to AAA2351 and AAA19223 to AAA2351 and AAA2350 to AAA23595 represent ribozyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       stability of an mRNA encoding an angiogenic factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel ribozymes for modulating the synthesis, expression and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC
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                                              subunit alpha-6, or integrin subunit
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Coeshott C,
                                                   beta-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          McSwiggen JA;
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Matches Query Match Local l Similarity
14; Conserv Conservative 0.9%; 0; Score 12.8; I Pred. No. 2.5e 0; Mismatches 2.5e+02; DB 1; Length 17; Indels 0, Gaps

0

17

B₽;

7 A,

4 C; 1 G;

5 U;

0 other;

RESULT 299 AAA18875 밁 AAA18875 standard; RNA; 17 17 TAATGITGGAAATCTT 2

2348 TAATGTGGGAGATCTT 2363

Human TIE-2 substrate sequence SEQ ID NO:2101. 19-JUN-2000

(first entry)

Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; anglogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antiporiatic; ARND; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome: ss.

Homo sapiens

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RESULT 300
AAA20440/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RNA Cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, and AAA17168 to AAA17560 and AAA17623 to AAA18484 represent their corresponding target sequences; AAA17683 to AAA18385 and AAA18386 to AAA18185 to AAA21856 to AAA21868 represent their corresponding target sequences; AAA18185 to AAA21868 represent their corresponding target sequences; AAA18185 to AAA2185 and AAA23363 to AAA23342 represent ribozyme sequence for integrin subunit beta 3, and AAA23476 co AAA23343 to AAA23343 to AAA23343 to AAA23343 to AAA23343 to AAA2385 to AAA23343 to AAA23475 and AAA23476 co AAA23343 to AAA23476 co A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA23422 represent their corresponding target sequences. The ribozymes the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, posiasis, verruca vulgaris, angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2,
                      Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antiporiatic; ARNU; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel ribozymes for modulating the synthesis, exprestability of an mRNA encoding an angiogenic factors
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                                                                                                                                                                                                                        Integrin alpha 6 subunit substrate sequence SEQ ID NO:3666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present invention describes enzymatic nucleic acid molecules with cleaving activity, which specifically cleave RNA encoded by an arr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2198 TAGCAGACTTTGGACT 2213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PA,
degeneration; psoriasis; verruca vulgaris;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                          standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             uAGCAGAUUUUGGAUU 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 BP; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      subunit alpha-6, or integrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                              (first
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                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>ა</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 12.8;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Coeshott
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.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
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(first entry) σ

AAA20441 standard; RNA; 17

Integrin 19-JUN-2000

alpha

subunit substrate sequence

SEQ ID NO:3667

Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme;

뭉 Ş

Matches

Conservative

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Score 12.8; I Pred. No. 2.5e 0; Mismatches

2.5e+02; DB 1;

Length 17;

0

Gaps

0

0.9%;

2183 16

ACAATGTGATGAAAAT 2198

Best Local Similarity

Query Match

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The present invention describes enzymatic nucleic acid molecules with CC RNA cleaving activity, which specifically cleave RNA encoded by an aryl chydrocarbon nuclear transporter (ART) gene, an integrin subunit beta 3 CC gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, CC and AAA17168 to AAA17623 to AAA18385 and AAA19087 to CC corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086 and AAA19155 to AAA23322 represent their corresponding target sequences; CC AAA19223 to AAA23361 and AAA21501 to AAA21595 represent ribozyme sequences for integrin alpha 6 subunit, and AAA23362 to AAA21500 and AAA21596 to AAA21688 represent their corresponding target sequences; AAA21689 to AAA21689 to AAA237475 and AAA23342 represent ribozyme sequence for integrin subunit beta 3, and AAA237476 to AAA23343 to AAA23342 represent their corresponding target sequences. The ribozymes of CC AAA21422 represent their corresponding target sequences. The ribozymes of CC AAA21422 represent their corresponding target sequences. The ribozymes of CC AAA21422 represent their corresponding target sequences. The ribozymes of CC AAA21422 represent their corresponding target sequences. The ribozymes of CC AAA21422 represent their corresponding target sequences. The ribozymes of CC AAA21422 represent their corresponding target sequences. The ribozymes of CC AAA21422 represent their corresponding target sequences. The ribozymes of CC AAA21422 represent their corresponding target sequences. The ribozymes of CC AAA21422 represent their corresponding target sequences.
                                                                                                                                   stability of an mRNA encoding angiogenic factor, especially ARNT, integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgari angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome and other syndromes and diseases related to the levels of ARNT, Tie-2
Sequence
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Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 55; Page 145; 305pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors
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     17
                                                                                                subunit
BP;
5 A;
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                                                                                                alpha-6,
3 C; 1 G; 8 U; 0 other;
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                                                                                           integrin subunit
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                                                                                                     beta-3
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                                                                                                                                                                                                                                                                                                verruca vulgaris,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88
                                                                                                                                                                                                syndrome,
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RESULT 302
AAA20859
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                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA23422 represent their corresponding target sequences. The ribozymes of the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2,
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AAA20859 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-591315/50
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                                                                                                                                                                                                                                                       2181 AAACAATGTGATGAAA 2196
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                                                                                                                                                                                      17 ATACAATGTAATGAAA 2
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                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; 3 C; 1 G; 9
                                                                                                                                                                                                                                                                                                                                                                    0.9%;
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                                                                                                                                                                                                                                                                                                                                     <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                        Score 12.8; DB 1;
Pred. No. 2.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    other;
                                                                                                                                                                                                                                                                                                                         2.5e+02;
2;
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Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                         07-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; 88
                                                                                                                                                                                                                                                                                                                                                                                                 27-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                         Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                 98US-0079678.
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                                                                                                                                                                                                                                                                                                                                                         Jarvis T,
                                                                                                                                                                                                                                                                                                                                                         Coeshott C, McSwiggen
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55; Page 172; 305pp; English.

Claim

Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors ${\tt -}$

The present invention describes enzymatic nucleic acid molecules with CC RNA cleaving activity, which specifically cleave RNA encoded by an aryl CC hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 CC gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to CC AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, CC and AAA17168 to AAA17560 and AAA17685 to AAA17684 represent their CC corresponding target sequences; AAA17685 to AAA13885 and AAA19087 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA19086 to AAA19155 to AAA19222 represent their corresponding target sequences; AAA19233 to AAA19155 to AAA29361 and AAA21501 to AAA2155 represent ribozyme sequences; AAA19223 to AAA21688 represent their corresponding target sequences; AAA19223 to AAA21688 represent their corresponding target sequences; AAA19223 to AAA21688 represent their corresponding target sequences; AAA21889 to AAA22475 and AAA23263 to AAA23342 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA23262, AAA23343 to AAA23422 represent their corresponding target sequences. The ribozyme sof the invention are used for modulating the synthesis, expression and/or c stability of an mRNA encoding angiogenic factor, especially ARNT, CC integrin subunit beta 3, integrin subunit alpha 6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related CC macular degeneration (ARMD), inflammation, and arthritis, as well as concusar syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndrome, and disease related to the levels of ARNT, Tie-2, integrin subunit beta-3 are integrin subunit beta-3. subunit alpha-6, or integrin subunit beta-3.

밁 ફ Matches Query Match Best Local 2191 ATGAAAATAGCAGACT 2206 ш Similarity AAGAAAAUACCAGACU 16 Conservative 0.9%; 2 Score 12.8; Pred. No. 2. Mismatches 2.5e+02; ches 2; DB 1; Length 17; Indels 0 Gaps

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Sequence

17 BP; 9

A; 4 C; 2 G; 2 U; 0 other;

RESULT 303

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AAA20859;

19-JUN-2000

(first entry)

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AAA21301 standard; RNA; 17 BP.

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2 UAUACAUUCAACAGAG 17

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AAA23422 represent their corresponding target sequences. The ribozymes of the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARWT; integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriass; verruca vulgaris, angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARMT, Tie-2, integrin subunit alpha-6, or integrin subunit beta-3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16755 to AAA1767 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, and AAA17168 to AAA17623 to AAA17624 represent their corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086 and AAA19155 to AAA19222 represent their corresponding target sequences; AAA19223 to AAA20361 and AAA21501 to AAA21595 represent ribozyme sequences for integrin alpha 6 subunit, and AAA20362 to AAA21500 and AAA22596 to AAA21688 represent their corresponding target sequences; AAA21689 to AAA21675 and AAA23263 to AAA23342 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA23262, AAA23343 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors -
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Sequence 17 BP; 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-MAR-1998;
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    A; 4 C; 2 G; 4 U; 0 other;
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2309 TATACACTCATCAGAG 2324

Query Match Best Local Matches

Local

Similarity

0.9%; Score 12.8; DB 1; Length 17; 68.8%; Pred. No. 2.5e+02; tive 3; Mismatches 2; Indels

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Gaps

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Query Match

0.9%;

Score 12.8;

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                                                                                                           AAA23422 represent their corresponding target sequences. The ribozymes the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARNT, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2, and other syndromes and diseases related to the levels of ARNT, Tie-2,
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Sequence
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tuberous sclerosis; pot-wine stain; Sturge Weber syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 55; Page 199; 305pp; English.
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                                                                              subunit alpha-6,
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8 A;
3 C; 2 G; 4 U; 0 other;
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                                                                              or integrin subunit beta-3.
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RESULT 305
AAA23037/c
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The present invention describes enzymatic nucleic acid molecules with CC RNA cleaving activity, which specifically cleave RNA encoded by an aryl CC hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 CC gene, an integrin alpha 6 subunit gene, or a Tie-2 gene AAA16775 to CC AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, CC and AAA17168 to AAA17560 and AAA17623 to AAA17684 represent their CC corresponding target sequences; AAA17625 to AAA18385 and AAA19087 to CC AAA19154 represent ribozyme sequence for Tie-2, and AAA18386 to AAA19086 CC and AAA19155 to AAA21622 represent their corresponding target sequences; AAA19232 to AAA21681 represent their corresponding target sequences; CC AAA19232 to AAA21681 represent their corresponding target sequences; CC AAA19232 to AAA21688 represent their corresponding target sequences; CC AAA21696 to AAA21688 represent their corresponding target sequences; CC AAA21698 to AAA22475 and AAA23263 to AAA231595 represent ribozyme sequence for integrin subunit beta 3, and AAA23476 to AAA23362, AAA23343 to CAA23689 to AAA22475 and AAA23263 to AAA23142 represent ribozymes of CC AAA23422 represent their corresponding target sequences. The ribozymes of the invention are used for modulating the synthesis, expression and/or citegrin subunit beta 3, and AAA22476 to AAA23362, AAA23343 to CAA23681 to AAA23426 represent ribozymes of the invention are used for modulating the synthesis, expression and/or citegrin subunit beta 3, integrin subunit alpha 6, or Tie-2. They are cespecially used to treat cancer, diabetic retinopathy, age related CC macular degeneration (ARMD), inflammation, and arthritis, as well as correspondence of the invention of tuberous soleromis, pot-wine stains, Sturge Weber CC syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndrome, and diseases related to the levels of ARMT, Tie-2,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 54;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       305pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Coeshott C, McSwiggen JA;
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                                                                                                  RESULT 306
AAX16147/c
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Best Local
                                                                                                                                 The present invention describes neurofibromatosis type 2 (NF2) transcript isoforms. NF2 polynucleotides can be used for diagnosing NF2 diseases, for inhibiting growth of tumours associated with NF2 mutations (including expression from cDNA introduced in gene therapy vectors) and to raise antibodies (useful as tumour targeting agents, since specific isoforms are often tumour-specific) and as immunoassay reagents for detecting NF2-expression products. NF2 is a tumour suppressor protein, and so has anticancer activity. The present sequence represents a PCR primer for mouse NF2, from an example of
                                                                                              Sequence 17
                                                                                                                                                                                                                                                             Example; Column 14; 45pp; English
                                                                                                                                                                                                                                                                                    Proteins from neurofibromatosis type 2 transcript isoforms for diagnosis or inhibition of tumours, and generation of a
                                                                                                                                                                                                                                                                                                                                                  Bianchi
                                                                                                                                                                                                                                                                                                                                                                                                   10-JAN-1994;
04-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; neurofibromatosis type 2; NP2; tumour suppressor; cancer; diagnosis; gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse neurofibromatosis type 2 PCR primer 5m9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX16147
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                                                            Local
                        1579
                                                                                                                        present invention.
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                                                 l Similarity
14; Conser
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                       TCCTCCATGAACTCCA 1594
TTCTCCATGTACTCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA;
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                                                                                               BP; 6 A; 1 C; 7 G; 3 T; 0 other;
                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                  Kley NA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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96US-0628145.
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                                                           Score 12.8;
Pred. No. 2
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Pred. No. 2.
                                                  Mismatches
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                                              3; L-
2.5e+02;
2;
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2.5e+02;
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RESULT 307 AAV93558/c

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CC capable of modulating a process in a biological system. The method comprises: (a) introducing into the system a random library of nucleic comprises: (a) introducing into the system a random library of nucleic comprises: (a) introducing into the system a random library of nucleic comprises: (a) introducing into the system a random library of nucleic cacid catalysts (NAC) having a substrate binding domain (SBD), comprising carried and catalytic domain (CD); and (b) identifying NAC in systems where modulation has occurred and/or determining the sequence compared to the SBDs in such systems. Nucleic acid molecules caused to modulate gene expression in plant and mammalian cells and to cleave target nucleic acid, particularly for treating comprises caused by specific RNA, e.g. cancer, inflammation, systemic diseases caused by specific RNA, e.g. cancer, inflammation, comprises in non-hepatic ascites and infection. They may also be used to detect genetic drift and mutations in diseased cells and to determine corresponding to the Raf gene, are used to treat cancer, restenosis, concression of the Raf gene, are used to treat cancer, restenosis, concreases stability against nuclease and activity any condition associated increases stability against nuclease and activity. AAV90922 to AAV93877 compresent NACs that can be used in the method, specifically for modulating the expression of a Raf gene.
Matches
                                Query Match
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02-OCT-1997;
02-OCT-1997;
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09-MAY-1997;
09-JUN-1997;
03-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme; target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification; synthesis; deprotection; purification; cancer; inflammation; psoriasis; non-hapatic ascites; infection; genetic drift; restenosis; rheumatoid arthritis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 177; Page 170; 259pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Karpeisky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC
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                  Local
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14;
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                  Similarity
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Reynolds M,
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                                                                    BP; 5
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97US-0061321.
97US-0061324.
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97US-0046059.
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97US-0051718
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, Matulic-Adamic J, McSwiggen JA;
Sweedler D, Thompson J, Workman CT;
0;
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                                Score 12.8; DB 1;
                  Pred. No.
Mismatches
                                                                    U; 0 other;
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2.5e+02;
thes 2;
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09-MAY-1997;
09-JUN-1997;
03-JUL-1997;
03-JUG-1997;
02-AUG-1997;
02-OCT-1997;
02-OCT-1997;
                                                with endomulease activity and catalytic activity, from the present invention, are used to modulate gene expression in plant and mammalian cells and to cleave target nuclei cally particularly for treating systemic diseases caused by specific RNA, e.g. cancer, inflammation, psoriasis, non-hepatic ascites and infection. They may also be used to detect genetic drift and mutations in diseased cells and to determine c-raf RNA. Specifically NAC with RNA-cleaving activity that modulate expression of the Raf gene, are used to treat cancer, restenosis, psoriasis or rheumatoid arthritis, or generally any condition associated with the level of c-raf. Introduction of sugar/phosphate modifications
                                                                                                                                                                                                             A method has been developed for the identification of a nucleic acid capable of modulating a process in a biological system. The method comprises: (a) introducing into the system a random library of nucleic acid catalysts (NAC) having a substrate binding domain (SBD), comprising a random sequence, and a catalytic domain (CD); and (b) identifying NAC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules with endonuclease activity and catalytic activity. from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Beaudry A,
Karpeisky ,
Parry T, 1
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                                                                                                                                                                                                                                                                                                                                                                              Claim 177; Page 169; 259pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-009494/01.
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              increases stability against nuclease and activity. AAV90922 to AAV93877 represent NACs that can be used in the method, specifically for
                                                                                                                                                                                                                                                                                                                                                                                                                 triphosphates used as antiviral agents and synthons
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97US-0049002.
97US-0051718.
97US-0056808.
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97US-0061324.
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, Matulic-Adamic J, McSwiggen JA;
Sweedler D, Thompson J, Workman
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RESULT 309
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Best Local Simi
Matches 14;
A.method has been developed for the identification of a nucleic acid capable of modulating a process in a biological system. The method comprises: (a) introducing into the system a random library of nucleic acid catalysts (NAC) having a substrate binding domain (SBD), comprising a random sequence, and a catalytic domain (CD); and (b) identifying NAC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules with endonuclease activity and catalytic activity, from the present invention, are used to modulate gene expression in plant and mammalian cells and to cleave target nucleic acid, particularly for treating systemic diseases caused by specific RNA, e.g. cancer, inflammation, psoriasis, non-hepatic ascites and infection. They may also be used to determine
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09-MAY-1997;
09-JUN-1997;
09-JUL-1997;
03-JUL-1997;
22-AUG-1997;
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Parry T,
                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                      Claim 177; Page 146; 259pp;
                                                                                                                                                                                                                 triphosphates used as antiviral agents
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            target; substrate; catalyst;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human C-raf target site nucleotide position
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                                                                                                                                                                                                               ying new catalytic nucleic acid that modulates selected es - especially ribozymes that cleave Raf RNA for treating restenosis, and also new ribozymes and modified nucleoside phates used as antiviral agents and synthons
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                                                                                                                                                                                                                                                                                                                                                                   RIBOZYME PHARM INC.
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Reynolds M,
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Kisich K, Matulic-Adamic J,
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2.5e+02;
nes 2;
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, McSwiggen Ja;
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ion; Raf gene;
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Best Local S
Matches 9
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                           A method has been developed for the identification of a nucleic acid capable of modulating a process in a biological system. The method comprises: (a) introducing into the system a random library of nucleic acid catalysts (NAC) having a substrate binding domain (SBD), comprising a random sequence, and a catalytic domain (CD); and (b) identifying NAC
                                                                                                                                                                                                                                      Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons
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Reynolds M,
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Kisich K, Matulic-Adamic J, McSwiggen JA,
synolds M, Sweedler D, Thompson J, Workman
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AAV91069/c
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Claim 177; Page 148; 259pp; English
                               Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons
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Karpeisky A, Kisich K, Matulic-Adamic J, McSwiggen JA,
Parry T, Reynolds M, Sweedler D, Thompson J, Workman
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Pred. No. 2.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C. A method has been developed for the identification of a nucleic acid ccapable of modulating a process in a biological system. The method CC comprises: (a) introducing into the system a random library of nucleic CC acid catalysts (NAC) having a substrate binding domain (SBD), comprising CC a random sequence, and a catalytic domain (CD); and (b) identifying NAC CC in systems where modulation has occurred and/or determining the sequence CC of at least part of the SBDs in such systems. Nucleic acid molecules CC with endonuclease activity and catalytic activity, from the present CC invention, are used to modulate gene expression in plant and mammalian CC cells and to clease target nucleic acid, particularly for treating CC systemic diseases caused by specific RNA, e.g. cancer, inflammation, CC psoriasis, non-hepatic ascites and infection. They may also be used to detect genetic dirft and mutations in diseased cells and to determine CC -raf RNA. Specifically NACs with RNA-cleaving activity that modulate CC expression of the Raf gene, are used to treat cancer, restenosis, CC mith the level of c-raf. Introduction of sugar/phosphate modifications in creases stability against nuclease and activity. AAV9092z to AAV93877 CC represent NACs that can be used in the method, specifically for modulating the expression of a Raf gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                     09-MAY-1997;
09-JUN-1997;
03-JUL-1997;
22-AUG-1997;
02-OCT-1997;
02-OCT-1997;
                                                       Beaudry A, Beigelman
Karpeisky A, Kisich
Parry T, Reynolds M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; C-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme; target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification; synthesis; deprotection; purification; cancer; inflammation; psoriasis; non-hepatic ascites; infection; genetic drift; restenosis; rheumatoid arthritis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-MAY-1998;
                                                                                                                                                                                                                                                           05-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human C-raf target site nucleotide position 1505.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-FEB-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV91152 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 2 A; 9 C; 1 G; 5 U; 0 other;
                                                                                                                                                                                          (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                            Beigelman
                                                       sigelman L, Bellon L, Burgin A, Jarvis T;
Kisich K, Matulic-Adamic J, McSwiggen JA;
nolds M, Sweedler D, Thompson J, Workman
                                                                                                                                                                                                                                                                                     97US-0056808.
97US-0061321.
97US-0061324.
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97US-0046059.
97US-0049002.
97US-0051718.
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Pred. No. 2.5e+02
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 17;
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WPI; 1999-009494/01

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RESULT 313
AAV91151
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Best Local S
Matches 10
                                                             19-DEC-1997;
09-MAY-1997;
09-JUN-1997;
03-JUL-1997;
22-AUG-1997;
                                                                                                                                                                   05-MAY-1998;
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                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                    target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification; synthesis; deprotection; purification; cancer; inflammation; psortasis; non-hepatic ascites;
                                                                                                                                                                                                                                                                                                                                                           Human; c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme;
target; substrate; catalyst; modulation; expression; Raf gene;
                                                                                                                                                                                                                                                                                                                                                                                                               Human C-raf target site nucleotide position 1504.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV91151 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 177; Page 149; 259pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons
                                                                                                                                                                                                                                                                                                         infection; genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2484
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10; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGACTGTTGGCATGCA 2499
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
            97US-0049002.
97US-0051718.
97US-0056808.
97US-0061321.
97US-0061324.
97US-0064866.
                                                                                               97US-0068212.
97US-0046059.
97US-0049002.
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                                                                                                                                                                     98WO-US09249
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                                                                                                                                                                                                                                                                                                         drift; restenosis; rheumatoid arthritis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of a Raf gene.
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Pred. No. 2.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 17;
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SXCCCCCCCCCCCCCCCCX XX TTTTXX X TITXXX X TIXXXX X TIXXX X TIXXXX X TIXXX X TI
                                                                                                                     with endonuclease activity and catalytic activity, from the present invention, are used to modulate gene expression in plant and mammalian cells and to cleave target nucleic acid, particularly for treating systemic diseases caused by specific RNA, e.g. cancer, inflammation, psoriasis, non-hapatic ascites and infection. They may also be used to detect genetic drift and mutations in diseased cells and to determine c-raf RNA. Specifically NACs with RNA-cleaving activity that modulate expression of the Raf gene, are used to treat cancer, restenosis, psoriasis or rheumatoid arthritis, or generally any condition associated with the level of c-raf. Introduction of sugar/phosphate modifications increases stability against nuclease and activity. AAV99922 to AAV93877 represent NACs that can be used in the method, specifically for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      capable of modulating a process in a biological system. The method comprises: (a) introducing into the system a random library of nucleic acid catalysts (NAC) having a substrate binding domain (SBD), comprising a random sequence, and a catalytic domain (CD); and (b) identifying NAC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 177; Page 149; 259pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Beaudry A, Beigelman
Karpeisky A, Kisich
Parry T, Reynolds M,
                                                                                    modulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         triphosphates used as antiviral agents and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                has
17
                                                                               the expression
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, Kisich K, Matulic-Adamic J,
eynolds M, Sweedler D, Thompson
BP; 4 A; 3 C; 4 G; 6 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           been
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         developed for the identification of a nucleic ting a process in a biological system. The meth troducing into the system a random library of r
                                                                               e used in the method, of a Raf gene.
0 other;
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RESULT 314
AAF06118/c
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                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 10
                                                                                                        Ribozyme; erythropoietin;
                                                                                                                      Hammerhead
                                                                                                                                                                 AAF06118 standard; DNA; 17
                           12-APR-1999;
                                        11-APR-2000;
                                                       19-OCT-2000
                                                                                  Homo sapiens
                                                                                                 interferon alpha;
                                                                                                                                     16-FEB-2001
                                                                                                                                                                                                                   2484 GGACTGTTGGCATGCA 2499
                                                                                                                                                                                                                                  10;
                                                                                                                                                                                                      N
            RIBOZYME PHARM INC.
                                                                                                                                                                                                                                         Similarity
                                                                                                                      ribozyme substrate #2915.
                                                                                                                                                                                                                                   Conservative
                                         2000WO-US09721
                                                                                                                                     (first entry
                           99US-0129390.
                                                                                                  88.
                                                                                                                                                                                                                                        0.9%;
                                                                                                       granulocyte colony stimulating
                                                                                                                                                                                                      17
                                                                                                                                                                  ВP
                                                                                                                                                                                                                                 4.
                                                                                                                                                                                                                                        Score 12.8;
Pred. No. 2
                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                       2.5e+02
                                                                                                                                                                                                                                               DB 1;
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                                                                                                                                                                                                                                               Length 17;
                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                 Gaps
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Blatt L,

Zwick M,

Pavco P,

McSwiggen

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AAF07173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factors the production of erythropoietin, granulocyte colony stimulating factors are presented in the production of erythropoietin, granulocyte colony stimulating factors are presented in the production of erythropoietin, granulocyte colony stimulating factors are presented in the production of erythropoietin, granulocyte colony stimulating factors are presented in the production of erythropoietin, granulocyte colony stimulating factors are presented in the production of erythropoietin, granulocyte colony stimulating factors are presented in the production of expression of presented in the present
               The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of represeor genes encoding the TR2 Orphan receptor, ERA3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interferon alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoletin -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                                            Enzymatic a useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ribozyme; erythropoietin; granulocyte colony stimulating factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hammerhead ribozyme substrate #3430.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RIBOZYME PHARM INC
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                                                                                                                                                                                                                                                                                               and antisense nucleic acid inhibition of repressor genes, r producing e.g. granulocyte colony stimulating factor interferon alpha and erythropoietin -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zwick M,
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87.5%;
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2.5e+02;
ches 2;
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                                                                         The present invention describes nucleic acids (A) that interact stably consists with a target sequence and contain at least one phosphoro(di)thioate clink, having endonuclease activity. (A), and more generally any ccatalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or cc endometrium), in vivo or by transforming cells ex vivo and implanting cc endometrium), in vivo or by transforming cells ex vivo and implanting cc estrogen receptor. Because of the high selectivity for targeted RNA, (A) cc can also be used to correlate inhibition of gene expression with cc alterations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that creativity, and as research reagents (for RNA, in the same way that creativity and as research reagents (for RNA). The combination of condifications in (A) improves resistance to nucleases, binding affinity and/or activity. AAA23503 to AAA24747 represent oestrogen receptor corresponding target sequences. AAA25992 represent coestrogen coeresponding target sequences. AAA26219 to AAA26218 represent cother ribozyme sequences and AAA26219 to AAA26218 represent cother ribozyme sequences and antisense oligonucleotides used in the example first in of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New nucleic acids that interact,
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23-JUN-1998;
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14; Conserv
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Zwick M, Jarv
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is T, Woolf T,
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                                         Clikk, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or cendometrium), in vivo or by transforming cells ex vivo and implanting treated cells, or for other conditions associated with levels of cestrogen receptor. Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with called a leverations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that restriction endonucleases are used with DNA). The combination of therapeutic and/or activity. AAA3503 to AAA24748 to AAA25992 represent their corresponding target sequences, and AAA26105 represent their corresponding target sequences. AAA22619 to AAA26718 represent their corresponding target sequences. AAA22610 to AAA2671 represent their their corresponding target sequences. AAA22610 to AAA2671 represent their corresponding target sequences. AAA2610 to AAA2671 represent their corresponding target sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acids that interact, sequences, used to treat cancer
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                                                                                                                                                                                                                                                                                                                                                                present invention describes nucleic acids (A) that interact stably
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Jarvis T, Woolf T,
                                present invention.
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Haeberli
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Query Match Best Local S

Similarity

0.9%;

Score 12.8; DB 1; Pred. No. 2.5e+02;

Length 17;

Query Match Best Local Matches

14;

Similarity

0.9%;

Score 12.8; DB 1; Pred. No. 2.5e+02; Mismatches

DB 1;

Length 17; Indels

0

Gaps

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0 other;

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RESULT 318
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ID AAA255
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                                            can also be used to correlate inhibition of gene expression with alterations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that medification endonucleases are used with DNA). The combination of modifications in (A) improves resistance to nucleases, binding affinity and/or activity. AAA23503 to AAA24747 represent oestrogen receptor hammerhead ribozyme sequences, and AAA24748 to AAA25932 represent their corresponding target sequences. AAA2593 to AAA26105 represent oestrogen receptor harpin ribozyme sequences. AAA2593 to AAA26107 to AAA26118 represent their corresponding target sequences. AAA2593 to AAA2617 represent oestrogen receptor harpin ribozyme sequences. AAA2619 to AAA26218 represent their ribozyme sequences. AAA2619 to AAA26218 represent their corresponding target sequences. AAA2619 to AAA26217 represent their ribozyme sequences and antisense oligonucleotides used in the
                                                                                                                                                                                                                                                         The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(di)thioate link, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or endometrium), in vivo or by transforming cells ex vivo and implanting treated cells, or for other conditions associated with levels of
                                                                                                                                                                                                                                                                                                                                                                                                   Claim
Sequence 17
                                   exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acids that interact,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-013248/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Thompson JD,
Reynolds M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       anticancer; breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oestrogen receptor hammerhead ribozyme target sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                     sequences, used to treat cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-APR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                 77; Page 81; 148pp; English
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Zwick M, Jar
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98US-0103636.
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 A
                                   the present invention.
 3 C; 1 G; 8 T; 0 other;
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vis T, Woolf T,
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Haeberli P;
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TTTTGGTAACAGAAAA 2183

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RESULT 319
ABA78913
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27-MAR-2000;
01-JUN-2000;
30-OCT-2000;
                                             the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor V, Factor VII, Factor IX, haemoglobin alpha locus 1 (HBA1), haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6, apolipoprotein E (APOE), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGT1), amyloid precursor protein (APC), presenilin-1 (PSEN1) and presenilin-2 (PSEN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP, PSENI; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentlin-1; Alzheimer; disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification -
                               Alzhermer
various syndromes. r
                                                                                                                                                                                                                                                  The present invention provides single-stranded oligonucleotides which (be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at
                                                                                                                                                                                                                                                                                                                                              Claim 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-639230/73
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17
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; 2000US-192179P.
; 2000US-208538P.
; 2000US-244989P.
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1 C; 5 G; 5 T; 0 other;
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                                   invention.
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Query Match Best Local : Matches

0.9%; Similarity 87.5%;

14;

Conservative

0; Mismatches

Score 12.8; DB Pred. No. 2.5e+

2.5e+02;

Length 17; Indels

0;

Gaps

0

Best Local Similarity

Query Match

17

BP;

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5 ი 0.9**%;** 87.5%;

μ o O

6 T; 0 other;

Score 12.8; DB 1; Pred. No. 2.5e+02;

Length 17;

Sequence

BP; 6 A;

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RESULT 320
ABA78914/c
ID ABA7899
XX ABA7899
XX ABA7899
XX ABA789
XX ABA789
XX ABA789
XX APC mn
XX Human
XX Human
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                                                                                         The present invention provides single-stranded oligonucleotides which can to be used for the targeted alteration of genomic sequences, where the coligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, tretinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor V, Factor VII, Factor IX, haemoglobin alpha locus (CDKN2A), APC, Factor V, Factor VIII, Factor IX, haemoglobin alpha locus (CDKN2A), APC, HAPOE), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGTI), amyloid precursor protein (APC), presenilin-1 (PSEN1) and CC presenilin-2 (PSEN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemoghilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting
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27-MAR-2000;
01-JUN-2000;
30-OCT-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical modification -
                                                                   oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 7; Page 147; 294pp; English
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; 2000US-192179P.
; 2000US-208538P.
; 2000US-244989P.
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ABA80468
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                                                                                                                                                                                                               The present invention provides single-stranded oligonucleotides which can be used for the targeted alteration of genomic sequences, where the oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CTFR, cyclin-dependent kinase inhibitor 2A (CDKN2A), APC, Factor V, Factor VIII, Factor IX, haemoglobin alpha locus 1 (HBA1), haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6, apolipoprotein E (APOB), LDL receptor (LDLR), UDP-glucuromosyltransferase (UGTI), amyloid precursor protein (APC), presenilin-1 (PSEN1) and presenilin-2 (PSEN2). These can be used in the gene therapy of diseases enoth as cannon adenosine descriptory.
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                                                                                             such as cancer, adenosine deaminase deficiency, cystic fibrosis, haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-MAR-2000; 2000US-192176P.
27-MAR-2000; 2000US-192179P.
01-JUN-2000; 2000US-208538P.
30-OCT-2000; 2000US-244989P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABA80468 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYDE ) UNIV DELAWARE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  UDP-glucuronosyltransferase; amyloid precursor protein; presenilin-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EB,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        leotide for targeted alterations of genetic sequences and for cystic fibrosis, comprises at least one mismatch and chemical
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Query Match

Sequence 17

BP; 3 A; 3 C; 4 G; 7 T; 0 other;

Score 12.8; DB 1;

Length 17;

Sequence 17 BP; 7

A; 4 C;

ω <u>.</u>

3 T;

0 other

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RESULT 322
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                                                                                   The present invention provides single-stranded oligonucleotides which can CC be used for the targeted alteration of genomic sequences, where the CC oligonucleotide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at CC the following genes: adenosine deaminase, p53, beta-globin, cretinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A CC (CDKN2A), APC, Factor V, Factor VII, Factor IX, haemoglobin alpha locus 2 (HBA2), MLH1, MSH2, MSH6, apolipoprotein & RAPOB, LDL receptor (LDLR), UDP-glucuronosyltransferase CC (UGT1), amyloid precursor protein (APC), presenilin-1 (PSEN1) and CC presenilin-2 (PSEN2). These can be used in the gene therapy of diseases CC such as cancer, adenosine deaminase deficiency, cystic fibrosis, CC haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, CC Alzheimer's disease, melanoma, adenomatous polyposis of the colon and CC various syndromes. The present sequence is one of the gene correcting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
Matches 14; Conserv
                                 cyclin-dependent kinase inhibitor 2A; DKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UCPT; syndrome; APP; PSENI; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentlin-1; Alzheimer's disease; cytostatic; antisickling; antianaemic; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 7; Page 226; 294pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical
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01-JUN-2000;
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2000US-208538P.
2000US-244989P.
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Best Local :
                                                                                                                                                                      Matches
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Best Local
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                                                                                                                                                                                                                                             The present invention provides nucleic acid molecules capable of downregulating the expression of the human checkpoint kinase-1 (chk1) gene. These may be antisense or ribozyme sequences, and are useful in the treatment of diseases associated with conditions affected by Chk1 levels, including cancer. The present sequence is an oligonucleotide described in
                                                                                                                                                                                                                                                                                                                                              Novel nucleic acid molecule e.g., ribozymes or antisense nucleic acid molecules, which downregulates expression of a checkpoint kinase-1 gene, useful for treating colorectal, lung, breast or prostate cancer:
                                                                                                                                                                                                                                                                                                                                                                                            WPI;
 Human Chkl ribozyme substrate
                        09-OCT-2001
                                                                    AAH95254 standard;
                                                                                                                                                                                                               Sequence 17 BP; 6 A; 4 C; 2 G; 5 U; 0 other;
                                                                                                                                                                                                                                                                                                                 Claim
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RNA cleavage;
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                                                                                                                                                                     1 Similarity
10; Conserv
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FATTAEY A R.
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                                                                                                                                                                     Conservative
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                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             kinase-1; Chk1; antisense; ribozyme; gene therapy;
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                        entry)
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                                                                    ВÞ
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                                                                                                                                                                               Score 12.8; DB Pred. No. 2.5e+(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12.8; DB 1;
Pred. No. 2.5e+02;
0; Mismatches 2;
                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                 Booher RN,
                                                                                                                                                                              2.5e+02
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                                                                                                                                                                                                                                                                                                                                                  cancers
                                                                                                                                                                   Gaps
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SEQ ID NO:

Fattaey

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Jarvis T,

McSwiggen

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Booher RN,

Holman

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RESULT 325
AAH95836
ID AAH958
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                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides nucleic acid molecules capable of downregulating the expression of the human checkpoint kinase-1 (Chk1) gene. These may be antisense or ribozyme sequences, and are useful in the treatment of diseases associated with conditions affected by Chk1 levels, including cancer. The present sequence is an oligonucleotide described in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel nucleic acid molecule e.g., ribozymes or antisense nucleic acid molecules, which downregulates expression of a checkpoint kinase-1 gene, useful for treating colorectal, lung, breast or prostate cancers
                                                                                                                                                                                                                             09-OCT-2001 (first entry)
                                                                                                                                                                                                                                                    AAH95836;
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 6 A; 2 C; 0 G; 9 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; checkpoint kinase-1; Chkl; antisense; ribozyme; gene therapy; RNA cleavage; cancer; ss.
                                                      03-FEB-2000; 2000US-0179983
                                                                                                    09-AUG-2001.
                                                                                                                           WO200157206-A2.
                                                                                                                                                Homo sapiens.
                                                                                                                                                                      RNA cleavage;
                                                                                                                                                                                                        Human Chkl ribozyme substrate SEQ
                                                                                                                                                                                                                                                                          AAH95836 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; Page 66; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-496922/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fattaey AR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-FEB-2000; 2000US-0179983
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                                                                              02-FEB-2001; 2001WO-US03504
                                                                                                                                                                                 Human; checkpoint kinase-1; Chk1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                2163 AAATGTTTTGGTAACA 2178
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                                                                                                                                                                                                                                                                                                                                                                              l Similarity
                      RIBOZYME PHARM INC
FATTAEY A R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FATTAEY A R.
                                                                                                                                                                                                                                                                                                                                  AAATGTTTTGATAAAA 2
                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                        cancer;
                                                                                                                                                                                                                                                                          RNA;
                                                                                                                                                                                                                                                                                                                                                                                       0.9%;
                                                                                                                                                                        88.
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                                                                                                                                                                                                                                                                            ВP
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                                                                                                                                                                                                                                                                                                                                                                              Score 12.8; DB 1;
Pred. No. 2.5e+02;
0; Mismatches 2
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                                                                                                                                                                                 antisense; ribozyme; gene therapy;
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                                                                                                                                                                                                         NO: 1261.
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                                                                                                                                                                                                                                                                                                                                                                                                  Length 17;
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RESULT 326
ABK00843/c
ID ABK008
XX ABK008
XX ABK008
XX Human
XX Human;
KW Cerebr
KW DNAzym
KW B-cell
KW Inflam
KW Creutz
XX Homo 8
OS Synthe
PN W02001
XX W02001
XX W02001
XX W02001
XX Homo 8
OS Synthe
PN W02001
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Best Local
                                                                                                                                                                                                                                                                                              11-FEB-2000;
28-FEB-2000;
06-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; NNAZyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; NCL; immunocytoma; INC, immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        downregulating the expression of the human checkpoint kinase-1 (Chk1) gene. These may be antisense or ribozyme sequences, and are useful in the treatment of diseases associated with conditions affected by Chk1 levels, including cancer. The present sequence is an oligonucleotide described in the exemplification of the invention.
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                              WPI; 2001-607195/69
                                                                                                                                                                            (RIBO-)
(BLAT/)
(MCSW/)
                                                                                                                                                                                                                                                                                                                                                                                                                   09-FEB-2001; 2001WO-US04273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Creutzieldt-Jakob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chemotherapy-induced neuropathy; amyotrophic lateral sclerosis;
Parkinson's disease; ataxia; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human NOGO Inozyme #113.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK00843 standard; RNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-496922/54.
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                                                                                                                                                                                                        RIBOZYME PHARM
BLATT L.
                                                                                                                                             CHOWRIRA B M.
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                                                                                                                                                                            MCSWIGGEN J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAUCUCAGACUUUGG 16
                                                                                   McSwiggen J,
                                                                                                                                                                                                                                                                                              ; 2000US-181797P.
; 2000US-185516P.
; 2000US-187128P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disease; muscular dystrophy; neurodegenerative disease
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                                                                                         Chowrira
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Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neuritogrowth inhibitor gene useful for treating, e.g., lymphoma, leukemia and central nervous system injury or neurite , leukemia,

79; 200pp; English

The invention relates to a nucleic acid molecule which down regulates cexpression of a CD20 gene and a nucleic acid molecule which down CC regulates expression of a neurite growth inhibitor gene (NGCD).

CC The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CC DNAzyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a CC DNAzyme) an Inozyme (an endolytic nucleic acid cleaving RNA with a NYN molecule (cleaving RNA with a YGY motif). The CD20-targetting nucleic acid is used CC (cleaving RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is correctly pranticed with a cell to reduce CC preferably Mg^2+. Furthermore, it may be contacted with a cell to reduce CC associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting complete cacid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular NHL, lymphocytic leukaemia, B-cell munocytoma (IMC), small B-cell lymphocytic leukaemia, B-cell munocytoma (IMC), small B-cell lymphocytic lymphoma, immune CC immunocytoma (IMC), small B-cell lymphocytic lymphoma, immune CC divalent cation that is preferably Mg^2+. Furthermore, the nucleic acid may be contacted with a cell to reduce NNA of the NOGO gene in the presence of a CC divalent cation that is preferably Mg^2+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the cell and CC may be contacted with a cell to reduce NOGO activity of the cell and CC treat a patient having a condition associated with the level of NOGO. The treat a patient having a condition associated with the level of NOGO. The central nervous system (CNS) injury and cerebrovascular accident (CVA, chemother), allowed the selections (MS), chemother, allowed the selections (MS), and/or other neurodegenerative disease disease, muscular dystrophy, and/or other neurodegenerative states which respond to the modulation of NOGO expression. T present sequence is an inozyme of the invention. disease

17 BP; 1 A; 8 C; 7 G; 1 U; 0 other;

Matches Query Match Local 1651 CTGGCAGGGGTCTCCG 1666 l Similarity 14; Conser Conservative 0.9%; 87.5%; 0; Score 12.8; Pred. No. 2. Mismatches 2.5e+02; DB 1; Length 17; Indels 0 0

밁 ABK00845 standard; RNA; 17 17 CCGGCAGGGGTCCCCG N

Human NOGO Inozyme #115

(first entry

RESULT 327
ABKO0845/c
ID ABKO08
XX ABKO08
XX II 2-MAR
XX II 2-MAR
XX Human
XX Horab
XX Human
XX Human Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic, neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NCG0; hammerhead ribozyme; muscular; CD20; neurite growth inhibitor gene; NCG0; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; be-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; NCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovasacular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkinson's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease

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                                                                                                                                                                                                                                                        CC DNazyme) an inozyme (an endolytic nucleic actid cleaving an RNA molecule consessing an NCH motif), a G-Cleaver (cleaving RNA with a NRN molecule content) pyr an amberzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with a rinzyme (cleaving RNA with a rinzyme) (cleave RNA of CD20 in the presence of a divalent cation that is contacted with the level of cD20. The treatment may further comprise the CD20 activity of the cell and treat a patient having a condition consociated with the level of CD20. The treatment may further comprise the cuse of one or more therapies. In particular, the CD20 targetting conjuncted acid may be used to treat lymphoma, lewhaemia, B-cell vow-grade or follicular NHL, lymphocytic leukaemia, HIV (human conjuncytoma (IMC), small B-cell lymphocytic leukaemia, HIV (human conjuncytoma (IMC), small B-cell lymphocytic lymphoma, immune conjuncted with a cell to cleave RNA of the NOGO gene in the presence of a conjuncted with a cell to reduce NOGO activity of the cell and creat a patient having a condition associated with the level of NOGO. The creat a patient having a condition associated with the level of NOGO. The central nervous system (CNS) injury and cerebrovascular accident (CVA, chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), character which respond to the modulation of NOGO expression. The contacted with respond to the modulation of the invention.
                                                                                                                          Matches
                                                                                                                                                        Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-FEB-2000;
28-FEB-2000;
06-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGG0). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or the nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or the nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or the nu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
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Synthetic.
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(BLAT/)
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                                                                                                                                                                                                                                                    Sequence 17
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                                                           1650 GCTGGCAGGGGTCTCC 1665
16
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BLATT L.
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                                                                                                                                                           Similarity
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GCCGGCAGGGGTCCCC
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; 2000US-185516P.
; 2000US-187128P.
                                                                                                                                                                                                                                                    BP; 1 A; 8 C; 7 G; 1 U; 0 other;
                                                                                                                          Conservative
                                                                                                                                                    0.9%;
87.5%;
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                                                                                                                                                           Score 12.8; DB 1;
Pred. No. 2.5e+02;
                                                                                                                          Mismatches
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                                                                                                                                                        2.5e+02;
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RESULT 328
ABK02092/C
ID ABK020
XX Human
XX Human
XX MCL; 1
KW MUSCL; 1
KW MCL; 1
KW M
The invention relates to a nucleic acid molecule which down regulates CC expression of a CD20 gene and a nucleic acid molecule which down CC regulates expression of a neurite growth inhibitor gene (NOGO).

CC The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CC DNazyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a CC DNazyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN CC (cleaving RNA with a CP1). The CD20-targetting nucleic acid is used cc (cleave RNA of CD20 in the presence of a divalent cation that is cc preferably Mg^2+. Furthermore, it may be contacted with a cell to reduce CC CD20 activity of the cell and treat a patient having a condition cassociated with the level of CD20. The treatment may further comprise the CC use of one or more therapies. In particular, the CD20 targetting cc lymphoma, low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky CC low-grade or follicular NHL, lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-Hodgkin's lymphoma (MCL), cimmunodeficiency virus) associated NHL, mantle-cell lymphoma (MCL), small B-cell lymphocytic leukaemia, immune cc immunocytopaemia, and inflammatory arthropathy. The NoGO-targetting cc nucleic acid is used to cleave RNA of the NoGO gene in the presence of a collect of a cleave RNA of the NoGO gene in the presence of a collect of a cleave RNA of the NoGO gene in the presence of a collect of a cleave RNA of the NoGO gene in the presence of a collect of a cleave RNA of the NoGO gene in the presence of a cleave RNA of the NoGO gene in the presence of a cleave RNA of the NoGO gene in the presence of a cleave RNA of the NoGO gene in the presence of a cleave RNA of the NGGO gene in the cleave and cleave RNA of the NGGO gene in the collection of a cleave RNA of the NGGO gene in the cleave cleave RNA of the NGGO ge
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28-FEB-2000;
06-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-)
(BLAT/)
(MCSW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
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; 2000US-187128P.
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RESULT 329
ABK02103/c
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                                                                                                                                                                                                                                  11-FEB-2000;
28-FEB-2000;
06-MAR-2000;
constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system inimm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      divalent cation that is preferably Mg^2^+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to treat central nervous system (CNS) injury and cerebrovascular accident (CNA, stroke). Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), parkinson's disease, ataxia, Huntington's disease, Creutzfeldt-Jakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present sequence is a DNAzyme molecule of the invention.
                                                                                                                                                                     (RIBO-)
(BLAT/)
(MCSW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkinson's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NO20; hammerhead ribozyme; DNAzyme; infozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia;
                                                                                                                                                                                                                                                                                                       09-FEB-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
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                                                                                                                                                                RIBOZYME PHARM
BLATT L.
MCSWIGGEN J.
                                                                                                                                                   CHOWRIRA B M.
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                                                                                                                McSwiggen J,
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; 2000US-185516P.
; 2000US-187128P.
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                                                                                                                                                                                                                                                                                                       2001WO-US04273.
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Pred. No. 2.5e+02;
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO).

CC The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CDNAzyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a CDNAzyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a CDAX, and the property of the angle of continuous continu
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          17
     BP; 1
A, 10 C;
     5 G;
     1 U;
     0 other;
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문 5 Matches Query Match Best Local 1653 GGCAGGGGTCTCCGAG 1668 17 14; Similarity Conservative 0.9%; N 0; Score 12.8; Pred. No. 2 Mismatches 2.5e+02 DB 1; 2; Length 17; 0; Gaps

0,

12-MAR-2002 (first entry) ABK02341 standard; RNA; 17 ВP

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; noctropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOG0; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; be-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunocytoma; INC; immunocytoma; IN

Homo sapiens Synthetic.

WO200159103-A2

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The invention relates to a nucleic acid molecule which down regulates compression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO).

CC The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CDNAzyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a CDNAzyme) an Inozyme (an endolytic nucleic acid cleaving RNA with a NYN molecule colis (cleaving RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is corrected with a cell and treat a patient having a condition colleave RNA of the cell and treat a patient having a condition colleave romagnetic formular treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky colleaved of one or more therapies. In particular, the CD20 targetting immunocytoma (NMC), small B-cell lymphoma, low-grade or follicular NHL, lymphocytic leukaemia, B-cell lymphoma, comparade or follicular NHL, lymphocytic leukaemia, B-cell lymphoma (NGC).

CC low-grade or follicular NHL, lymphocytic lymphoma (NGL), colleave RNA of the NGCO gene in the presence of a colleave romagnetic leukaemia, B-cell lymphoma (NGC).

CC divergrade or follicular NHL, lymphocytic lymphoma, immune collectic acid is used to cleave RNA of the NGCO gene in the presence of a collection of that is preferably Mg^2+. Furthermore, the nucleic acid can be contacted with a cell to reduce NGCO activity of the cell and treat a patient having a condition associated with the level of NGCO. The creat a patient having a condition associated with the level of NGCO. The central nervous system (CNS) injury and cerebrovascular accident (CVA, chick), Alzheimer's disease, dementia, multiple sclerosis (ALS), and insease, muscular dystrophy, and/or other neurodegenerative disease
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                                                                                                                         Matches
                                                                Query Match
Best Local :
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28-FEB-2000;
06-MAR-2000;
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(BLAT/)
(MCSW/)
                                                                                                                         Sequence
                                                                                                                                                                               states which
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                                                                  Similarity
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                                                                                                                                                                          muscular dystrophy, and/or other neurodegenerative hich respond to the modulation of NGCO expression. T
                                                                                                                         17
       GTCCAGCTCCTCCATG
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87.5%;
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         1587
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                                                                Score 12.8;
Pred. No. 2.
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                                              Mismatches
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le or neurite
leukemia,
                                              0
                                              Gaps
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RESULT 331 ABK02926/c ID ABK029

ABK02926 standard; RNA; 17

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17

N

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The invention relates to a nucleic acid molecule which down regulates CC expression of a CD20 gene and a nucleic acid molecule which down CC regulates expression of a neurite growth inhibitor gene (NGO).

CC The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CDNaxyme) an Inozyme (an endolyric nucleic acids (e.g. a ribozyme or a CDNaxyme) an ANCH motif), a G-cleaver (cleaving RNA with a NGN triplet), a zinzyme (cleaving RNA with a CP10 triplet), a zinzyme (cleaving RNA with a NGN triplet), a zinzyme (cleaving RNA with a CP20 in the presence of a divalent cation that is collary triplet), of the cell and treat a patient having a condition cassociated with the level of CD20. The treatment may further comprise the cuse of one or more therapies. In particular, the CD20 targetting cleaves of collicular NHL, lymphocytic leukaemia, B-cell lymphoma (NHL), bulky cleaving triplet acid is used to treat lymphoma, leukaemia, B-cell low-grade or follicular NHL, lymphocytic leukaemia, HIV (human collicit acid is used to cleave RNA of the NGO gene in the presence of a divalent cation that is preferably Mg^2+. Furthermore, the nucleic acid cold treat a patient having a condition associated with the level of NGO. The treatment may further comprise the use of one or more therapies.
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28-FEB-2000;
06-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RIBO-)
(BLAT/)
(MCSW/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 30; Page 143; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-FEB-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sapiens.
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BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CHOWRIRA B M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MCSWIGGEN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          McSwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; 2000US-181797P.
; 2000US-185516P.
; 2000US-187128P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001WO-US04273.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NGG0; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; he-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; inmunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; Takase creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   In particular, the NOGO-targetting nucleic acid may be used to treat central nervous system (CNS) injury and cerebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), parkinson's disease, ataxia, Huntington's disease, Creutzfeldt-Jakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The
                                                                         Nucleic acid molecules, e.g., enzymatic nucleic acids and constructs, which down regulate expression of a CD20 gene growth inhibitor gene useful for treating, e.g., lymphoma,
                                                                                                                                                                                                                               (RIBO-)
(BLAT/)
(MCSW/)
                             Claim 30; Page 149; 200pp; English
                                                             and central nervous system
                                                                                                                                                WPI; 2001-607195/69
                                                                                                                                                                               Blatt L,
                                                                                                                                                                                                                                                                                               11-FEB-2000; 2000US-181797P
28-FEB-2000; 2000US-185516P
06-MAR-2000; 2000US-187128P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human CD20 Inozyme #222.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK03271;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK03271 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 2 A; 7 C; 0 G; 8 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present sequence is a hammerhead ribozyme of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1362 TGGAAGAGAAAAGGAG 1377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14;
                                                                                                                                                                                                                                             RIBOZYME PHARM INC.
BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                CHOWRIRA B M.
                                                                                                                                                                                                                                 MCSWIGGEN J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TGTAAGAGAGAAGGAG 1
                                                                                                                                                                               McSwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.9%;
                                                                                                                                                                               Chowrira BM;
                                                             injury
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 12.8; DB 1;
Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                             d antisense
or neurite
a, leukemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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The invention relates to a nucleic acid molecule which down regulates

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RESULT 333
ABK03295/c
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cc The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a cc DNAzyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a conservation of a neurite growth inhibitor gene (NOGO)

cc possessing an NCH motif), a G-cleaver (cleaving RNA with a NYA molecule content of the presence of a divalent cation that is concerned to cleave RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is correctably Mg^2^+, Furthermore, it may be contacted with a cell to reduce CC CD20 activity of the cell and treat a patient having a condition consecution of concerned the level of CD20. The treatment may further comprise the cuse of one or more therapies. In particular, the CD20 targetting cc low-grade or follicular NHL, lymphocytic leukaemia, B-cell immunodeficiency virus) associated WHL, mantle-cell lymphoma (NHL), cc immunocytoma (IMC), small B-cell ymphocytic leukaemia, HIV (human concerned cacid is used to cleave RNA of the NGOO gene in the presence of a divalent cation that is preferably Mg^2^+. Furthermore, the nucleic acid may be contacted with a cell to reduce NGOO activity of the cell and creat a patient having a condition associated with the level of NGCO. The treat a patient having a condition associated with the level of NGCO. The creat a patient having a condition associated with the level of NGCO. The creat a patient having a condition associated with the level of NGCO. The creat a patient having a disease, dementia, multiple selerosis (NS), chemotherapy-induced neuropathy, amyocrophic lateral sclerosis (ALS), chemotherapy-induced neuropathy and content neuropassion. The present sequence is an inozyme of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                           Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; INC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis;
                                                                                                                                                                                                                   chemotherapy-induced neuropathy; amyotrophic lateral sclerosis;
Parkinson's disease; ataxia; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human CD20 Inozyme #246.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 6 A; 4 C; 4 G; 3 U; 0 other;
                                                                                                                                              Homo
                                                                                                                                                                                              Creutzfeldt-Jakob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK03295 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1703 CAAGAGATAAGCTGAC 1718
                                                                                                                                           sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%;
Similarity 81.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CAAGAGACAUGCUGAC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                              disease;
                                                                                                                                                                                              muscular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 12.8; DB 1;
Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                            neurodegenerative disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                      The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down completes expression of a neurite growth inhibitor gene (NOCO).

CC The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CDNAzyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA molecule CC possessing an NCH motif), a C-cleaver (cleaving RNA with a NYM control of the control of 
                                                                                                    RESULT 334
ABV78898
                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                            Query Match
Best Local
03-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RIBO-)
(BLAT/)
(MCSW/)
                                         ABV78898;
                                                                             ABV78898
                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 30; Page 149; 200pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-FEB-2000;
28-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-FEB-2001; 2001WO-US04273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CHOW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            central nervous system injury
                                                                                                                                                                                                                   1362 TGGAAGAGAAAAGGAG 1377
                                                                                                                                                                                17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RIBOZYME PHARM :
) BLATT L.
) MCSWIGGEN J.
                                                                                                                                                                                                                                                                          0.9%;
Similarity 87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CHOWRIRA B M.
                                                                                                                                                                                                                                                                                                                                      17
                                                                             standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               McSwiggen J,
                                                                                                                                                                                                                                                                                                                                      BP; 2
                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000US-181797P.
2000US-185516P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000US-187128P
(first entry)
                                                                                                                                                                                                                                                                                                                                      A; 7 C; 1 G; 7 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chowrira BM;
                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                            Score 12.8; DB 1;
Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                               Length 17;
                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                        Gaps
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RESULT 335 ABV78900 ID ABV789 XX AC ABV789

ABV78900 standard; DNA;

17 ₽₽

ABV78900

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                                                                                                                                   The present invention relates to human testis expressed Patched like CC protein (HTPL, see ABV78759 to ABBV8520). HTPL CC has two isoforms, with a few single base pair differences between the CC two. One of the single base pair changes introduces a premature stop CC codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL CC shares an overall structure organisation with the Patched protein. The CC shared structural features strongly imply that HTPL plays a role similar CC important in regulating male germ cell development, and the HTPL gene was CC mapped to human chromosome 10pl2-1. HTPL and its coding sequence are CC useful for diagnosing a disorder caused by mutation in HTPL, and in CC therapy and manufacture of a medicament for treatment or prevention of CC therapy and manufacture of a medicament for treatment or prevention of CC such disorder associated with decreased expression or activity of human CC foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, CC skeletal muscle or colon function. HTPL proteins and nucleic acids are CC clinically useful diagnostic markers and potenial therapeutic agents for CC manner from the invention. The present oligonucleotide was used in an CC example from the invention.
                                                                   Matches
                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JAN-2001;
30-JAN-2001;
23-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; halp germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel isolated human testis expressed Patched like protein useful for identifying agonist and antagonist and specific partners, and for treating subjects having defects in HTPL
                                                                                                                                   Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Page 82; 718pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-676582/73
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                                1518
                                                                 14;
2 GCCCAAGCTCACCAAA 17
                                                                                                                                                                     from
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                                GCACAAGCTGACCAAA 1533
                                                                                                                                   BP; 7
                                                                 Conservative
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2001WO-US00669.
2001US-0864761.
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2001WO-US00664.
2001WO-US00665.
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                                                                                                                                                                     invention.
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                                                                                0.9%;
                                                                                                                                   7 C; 2 G; 1 T; 0 other;
                                                                 0;
                                                                                Score 12.8; DB 1
Pred. No. 2.5e+02
                                                                   Mismatches
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                                                                                                   DB 1;
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                                                                                                  Length 17;
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                                                                   Gaps
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                                                                                                                                                                                              Matches
                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                            codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shared attructural structure organisation with the Patched protein. The shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL is mapped to human chromosome 10pl2.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an example for the structure of the present oligonucleotide was used in an example for the structure of the present oligonucleotide was used in an example for the structure of the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example for the present oligonucleotide was used in an example of the protein and the protein and the protein and the protein and th
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                                                                                                                                                                                              Sequence 17 BP; 6 A; 9 C; 1 G; 1 T; 0 other;
                                                                                                                                                                                                                                               example
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30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-MAY-2001;
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                            1519 CACAAGCTGACCAAAC 1534
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                                                                                               14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AEOMICA INC
                                                                                                                      Similarity
CCCAAGCTCACCAAAC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001WO-US00663.
2001WO-US00664.
2001WO-US00665.
2001WO-US00667.
2001WO-US00669.
2001WO-US00669.
2001WO-US00669.
                                                                                               Conservative
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                                                                                                                                                                                                                                               the invention.
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                                                                                                                   0.9%;
                                                                                            0;
                                                                                            Score 12.8; DB
Pred. No. 2.5e+1
0; Mismatches
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                                                                                                                 .5e+02;
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                                                                                                                                        Length 17;
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binding
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                                                                                            Gaps
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RESULT 336 ABV80429/c ID ABV804

ABV80429 standard; DNA; 17

RESULT 337

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                                                                                                                                                                                            The present invention relates to human testis expressed Patched like protein (HTPL, see ABY78759 to ABV78762 and ABB98519 to ABB98520). HTPL CC. has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stopy codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL CS shares an overall structure organisation with the Patched protein. The Start of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was important in regulating male germ cell development, and the HTPL gene was compared to human chromosome 10p12.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in the HTPL gene was such disorder associated with decreased expression or activity of human CC such disorders include disorders of testis, or adrenal, adult and CC foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and optenial therapeutic agents for male inferritity and cancer the proteins and nucleic acids are
                                                               Matches
                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL -
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                                                                                                                                 Sequence
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30-JAN-2001;
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30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                              2187 TGTGATGAAAATAGCA 2202
                                                                                                                                                                   infertility and cancer. ple from the invention.
17
                                                               14;
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                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AEOMICA
                                                                                                                                 17 BP; 5 A; 2 C; 1 G; 9 T; 0 other;
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2001WO-US00667.
2001WO-US00668.
2001WO-US00669.
2001US-0864761.
2001US-0327898.
                                                               Conservative
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2001WO-US00664.
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                                                                              0.9%;
                                                               o
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                                                                               Score 12.8;
Pred. No. 2.
                                                                                                                                                                                   The present oligonucleotide
                                                               Mismatches
                                                                                 2.5e+02;
                                                                                                 B
                                                                                               Length 17;
                                                               Indels
                                                                                                                                                                                     was used in an
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ABV80431/c

IID ABV80431/c

AC ABV804

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PR 30-JAN

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                                                                                                                                               Query Match
Best Local
                                                                                                                         Matches
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30-JAN-2001;
30-JAN-2001;
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                                                                                                                                                                                                                                           Sequence 17
                                                                                                                                                                                                                                                                                                        example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL -
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16
                                                                                                                                               Similarity
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ATGTTATAAAAATAGC
                                                         ATGTGATGAAAATAGC 2201
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2001WO-US00665.
2001WO-US00667.
2001WO-US00668.
2001WO-US00669.
                                                                                                                                                                                                                                           BP; 5
                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                        the invention.
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                                                                                                                                                                                                                                           A; 3 C; 1 G; 8
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                                                                                                                                               0.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English.
                                                                                                               Score 12.8; DB 1;
Pred. No. 2.5e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                           T; 0 other;
                                                                                                                                                                            Length 17;
                                                                                                                     Indels
                                                                                                                  0;
                                                                                                                  Gaps
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cc protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino (CC acids (S1, ABB83999), a sequence having 65% sequence identity to (S1), (S1) having 95% deviations, especially conservative substitutions or a CC fragment of the sequences comprising at least 8 contiguous amino acids. CC Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as CC downstream components of the signal transduction pathway. (I) is useful concoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL 1 including diagnosing and CC useful in gene therapy. (II) is useful for constructing microarrays which CC are useful for measuring and for surveying gene expression and creating cc are useful for measuring and for surveying gene expression and creating cc are useful for measuring and for surveying gene expression and creating creating transgenic non-human animals capable of producing the proteins. The CC of the invention.

CC Note: The nresent
ARUSULT 338
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XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human SH3 domain (POSH)-like signalling protein 1 POSHL-1, useful for treating disorders associated with expression or activity of human POSHL1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABV89408 standard; DNA; 17
                                                                                            specification, but
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; SEQ ID NO 121; 60pp + Sequence Listing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-684061/74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shannon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-MAY-2001;
10-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-JAN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene;
Rho GTPase; signal transduction; gene expression; cancer; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human POSHL1 scanning oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABV89408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (AEOM-) AEOMICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         therapy;
                                                                                         present sequence did not form part of the tion, but is based on sequence information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001WO-US00666.
2001WO-US00667.
2001WO-US00668.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001WO-US00670.
2001US-0864761.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002EP-0001165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001US-0328205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001WO-US00669.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2001WO-US00664.
2001WO-US00665.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001WO-US00663.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               transgenic; ss.
                                                             Patent Office.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NO 121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English
                                                                                         printed supplied
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     | polypeptide, decreased
                                                                                               ç
                                                                                               Derwent
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Query Match Best Local Matches

Similarity

0.9%;

0

Score 12.8; DB 1; Pred. No. 2.5e+02; 0; Mismatches 2;

Length 17; Indels

0

Gaps

0

Sequence 17

B₽;

5

₽,

4 C; 5 G; 3 T; 0 other;

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RESULT 339
ABV89411/c
ID ABV8941/c
ID ABV894
XX ABV894
XX ABV894
XX ABV894
XX Human;
KW Human;
KW Rho GT
XX Rho GT
X
                                                                                                                                              CC acids (S1, ABB8399), a sequence having 65% sequence of 730 amino (C2) having 95% deviations, especially conservative substitutions or a CC fragment of the sequences comprising at least 8 contiguous amino acids. CC Human POSH1 is a proto-oncogene/oncogene product that functions as an CC adaptor protein that interacts with Rho family small GTPases as well as CC downstream components of the signal transduction pathway. (I) is useful CC for identifying a specific binding partner. (I) and nucleic acids (II) CC encoding (I) are useful for diagnosing, monitoring disease and treating CC caused by altered expression of human POSHLI including diagnosing and CC useful for measuring and for surveying gene expression and creating CC useful for measuring and for surveying gene expression and creating CC are useful for measuring and for surveying gene expression and creating creating energy tensor to the proteins. The CC present sequence is that of a scanning oligonucleotide useful in examnles CC of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ś
                                                           specification, but
by the Europa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel human SH3 domain (POSH)-like signalling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1 -
                                                                                                           Note: The present sequence did not form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; SEQ ID NO 124; 60pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-684061/74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EP1239051-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene therapy; transgenic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; POSHL 1; SH3 domain; POSH-like signalling |
Rho GTPase; signal transduction; gene expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human POSHL1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABV89411;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABV89411 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CGGCTGAGTCCAGCTC 1580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CIGCIGAGITCAGCIC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001WO-US00666.
2001WO-US00667.
2001WO-US00668.
2001WO-US00669.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001WO-US00663.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2002EP-0001165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 scanning oligonucleotide SEQ ID NO 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001US-0864761.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001WO-US00665.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001WO-US00664.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001WO-US00670.
                                                               Patent
                                                               is based on sequence 
ent Office.
                                                                                        part of the information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein 1; oncogene;
; cancer; vaccine;
                                                                                                                       printed
                                                                                        supplied to Derwent
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Sequence 17

BP;

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A; 4

C; 5 ဂ္ N

T; 0 other

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RESULT 341
ABN00661/c
ID ABN006
XX
AC ABN006
XX
AC ABN006
XX
DT 29-MAY
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ABN83025
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                                                                                                                                                                               Matches 14;
                                                                                                                                                                                           Best Local Similarity
                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
            29-MAY-2002
                                                                                                                                                                                                                                                                                                           The invention relates to a novel process for a manufacturer to obtain customer orders for custom-designed biochips in an automated process. The invention also includes an automated system and process for providing a fully automated process for the design, manufacture and analysis of data
                                                                                                                                                                                                                                                                                                                                                                                                     Automated process for custom-designed biochip design, comprises obtaining desired target sequences from customer, creating sequencent motif for an array and applying the motif to a surface for later detection .
                                       ABN00661;
                                                              ABN00661 standard; DNA; 17
                                                                                                                                                                                                                                 Sequence 17 BP; 4 A;
                                                                                                                                                                                                                                                                       example of an array that may be designed using
                                                                                                                                                                                                                                                                                    designed
                                                                                                                                                                                                                                                                                                                                                                             Example 5; Page 21; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-APR-2000;
22-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ataxia telangiectasia; probe; biochip; array; capture; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABN83025;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABN83025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-017664/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200180155-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ataxia telangiectasia locus 56594896-SNeg-t capture probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (COMB-) COMBIMATRIX CORP.
                                                                                                                                                                                                                                                                                    biological array devices. The sequence represents a capture probe igned in the invention for the "sample ataxia" set of targets, as an
                                                                                                                                                     2104 GCCAGAGGCATGGAGT 2119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1563 TTCGGCTGAGTCCAGC 1578
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16 TTCTGCTGAGTTCAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14;
                                                                                                                             μ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ₽₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA; 17
                                                                                                                             GGCAGAGGCATGCAGT 16
                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000US-198045P
2000US-252880P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001WO-US12750
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Quarles
                                                                                                                                                                                                                                                                                                                                                                                                                                           for custom-designed biochip design, comprises
                                                                                                                                                                                           0.9%;
                                                                                                                                                                                                                                 3 C; 8 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ghazvini S;
                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                           Score 12.8;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 12.8; DB 1;
Pred. No. 2.5e+02;
                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                            2.5e+02;
                                                                                                                                                                                                                                                                        the method of the
                                                                                                                                                                                                      Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 17;
                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                creating sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                    suitable
                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:653

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of hgDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hgDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hgDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hgDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption consation, as therapeutic supplement in patients having specific deficiency in hgDMLP-1 production, and in vaccines or for replacement therapy. The polynuclectide sequences encoding hgDMLP-1 may be used for diagnosing a disorder associated with the expression of hgDMLP-1, in particular heart and skeletal muscle disorders. hgDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-MAY-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ଧ
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Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a human genome-derived myosin-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human
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                                                                                                                                                                                screening of the hGDMLP-1 sequence in the exemplification of the present
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                                                        ftp.wipo.int/pub/published_pct_sequence.
                                                                                           The sequence data for this patent did not form part of the printed iffication, but was obtained in electronic format directly from WIPO
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C; 5
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G; 2 T; 0 other;
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Query Match Best Local S Matches 14

Similarity

0.9%;

14;

Conservative

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Score 12.8; DB Pred. No. 2.5e+(0; Mismatches

DB 1; 2.5e+02; les 2;

Length 17; Indels

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Gaps

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RESULT 342
ABNO0662/c
ID ABNO06
XX Human;
XW Human;
XW Muscle
XX Homo s
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XX Homo s
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PP 25-MAY
XX 26-MAY
PR 26-MAY
PR 21-SEE
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PR 04-OCT
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PR 3
CC protein 1 (hgDMLP-1) The protein and polynucleotide sequences of thgDMLP-1 can be used in gene therapy and vaccine production. The CC hgDMLP-1 nucleic acids can be used as probes to detect, characterise CC and quantify hgDMLP-1 nucleic acids in samples, as amplification CC substrates, to provide initial substrates for the recombinant engineering CC of hgDMLP-1 protein variants having desired phenotypic improvements, and CC of thgMLP-1 proteins to raise antibodies that specifically recognise CC hgDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hgDMLP proteins, as specific Diomolecule capture probes for surface-enhanced laser desorption CC ionisation, as therapeutic supplement in patients having specific CC diagnosing a disorder associated with the expression of hgDMLP-1, in cc particular heart and skeletal muscle disorders. hgDMLP-1 is localised to
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30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a human genome-derived myosin-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
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27-SEP-2000;
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heart disease;
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RESULT 343
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invention.

N.B. The sequence data f
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       protein 1
hGDMLP-1
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27-SEP-2000;
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It invention describes a human genome-derived myosin-like (hGDMLP-1). The protein and polynucleotide sequences of land be used in gene therapy and vaccine production. The nucleic acids can be used as probes to detect, charactering
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ilarity 87.5%;
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                                                                                                                  ID 1107;
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0; Mismatches
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Pred. No. 2.
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                                                                                                                English
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2.5e+02;
les 2;
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RESULT 344
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Best Local S
Matches 14
                                                                                                            30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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21-SEP-2000;
27-SEP-2000;
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30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                     Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; hear muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                  30-JAN-2001;
30-JAN-2001;
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                                        (AEOM-)
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                                                                   2001WO-US00669.
2001WO-US00670.
2001US-266860P.
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2001WO-US00666.
2001WO-US00667.
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   30-JAN-2001;
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                                                                                                                                                                                                                                                                                              06-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                muscle; myosin; chromosome 22; gene therapy; vaccine; heart skeletal muscle disorder; amplicon; screening; ss.
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                                                                                       04-OCT-2000;
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2000US-234687P.
2000US-236359P.
2000GB-0024263.
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                              2001WO-US00661.
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t directly from WIPO
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RESULT 346
ABNO1121
ID ABNO11
XX ABNO11
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XX 29-MAY
DT 29-MAY
XX Human
XX Human;
XW Human;
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                               Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                      Human GDMLP-1
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                                                                                                                                                        29-MAY-2002
                                                                                                                                                                                                                        ABN01121 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 4 A;
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                                                                                                                                                                                                                                                                                                                                            1730 CCCTGGGAGAAGGTTG 1745
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87.5%;
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Pred. No. 2.5e+02;
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                                                                                     hGDMLP-1; heart;
                                                                                                                      ID NO:1113.
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Homo sapiens

heart disease;

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                                                                                                                                                                                                           CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 nucleic acids can be used as probes to detect, characterise CC and quantify hGDMLP-1 nucleic acids in samples, as amplification CC substrates, to provide initial substrates for the recombinant engineering CC of hGDMLP-1 protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may CC be used as immunogens to raise antibodies that specifically recognise CC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific concentration and/or amount specifically of hGDMLP proteins, as specific concentration, as therapeutic supplement in patients having specific CC deficiency in hGDMLP-1 production, and in vaccines or for replacement CC diagnosing a disorder associated with the expression of hGDMLP-1, in CC particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the correction.
                                                             Best Local Similarity Matches 14; Conserv
                                                                               Query Match
Best Local
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21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                                                                                                                          Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human
                                                                                                                                                                        N.B. The sequence data for this patent did not form papecification, but was obtained in electronic format
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protein 1 ()
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                   1730 CCCTGGGAGAAGGTTG 1745
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CCCTGGGAGAAAGTGG 16
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                                                                                                                          BP; 4 A; 3 C; 8
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2000US-23559P.

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                                                              Conservative
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                                                             0;
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                                                                           Score 12.8; DB 1;
Pred. No. 2.5e+02;
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RESULT 347 ABN01183 ID ABN01183 standard; DNA; 17 BP.

Sequence 17 BP; 9

A; 1 C;

7 G; 0 U; 0 other

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CC and quantify hGDMLP-1 nucleic acids in samples, as amplification CC substrates, to provide initial substrates for the recombinant engineering CC of hGDMLP-1 protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may CC be used as immunogens to raise antibodies that specifically recognise CC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption CC ionisation, as therapeutic supplement in patients having specific CC deficiency in hGDMLP-1 production, and in vaccines or for replacement CC therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for CC diagnosing a disorder associated with the expression of hGDMLP-1; in CC chromosome 22. The present sequence represents an oligomer used in the correction of the hGDMLP-1 sequence in the exemplification of the present
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30-JAN-2001;
                                           N.B. The sequence specification, but
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                                                                                                                                                                                                                                                                                                                         protein 1 (hGDMLP-1). The protein and polynucleotide sequences o hGDMLP-1 can be used in gene therapy and vaccine production. The
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27-SEP-2000;
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                           ftp.wipo.int/pub/published_pct_sequence
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21-SEP-2000;
27-SEP-2000;
                                                                            The present invention describes a human genome-derived myosin-like protein 1 (hgDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering
         of hgDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hgDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hgDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specifically of hGDMLP proteins, as specifically of hGDMLP proteins.
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biomolecule capture probes for surface-enhanced
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; myosin; chromosome 22; gene therapy; vaccine; heart disease;
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Pred. No. 2.5e+02;
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Disclosure;

SEQ ID 1571; 214pp; English

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RESULT 349
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
                                           New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1 -
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30-JAN-2001;
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27-SEP-2000;
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30-JAN-2001
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Pred. No. 2.5e+02;
0; Mismatches 2
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cc and quantify hGDMLP-1 nucleic acids in samples, as amplification cubstrates, to provide initial substrates for the recombinant engineering cof hGDMLP-1 protein variants having desired phenotypic improvements, and cf for expressing the proteins. The hGDMLP-1 proteins or polypeptides may cc leaved as immunogens to raise antibodies that specifically recognise cc hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption constation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for cd dagnosing a disorder associated with the expression of hGDMLP-1, in concommendation and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the exceening of the hGDMLP-1 sequence in the exemplification of the present
Sequence 17 BP; 2 A; 7 C; 5 G; 3 T; 0 other;
                                                                                     at ftp.wipo.int/pub/published_pct_sequence.
                                                                                                                                            N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
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망 Ś Query Match 0.9%; Best Local Similarity 87.5%; Matches 14; Conservative 1503 CAGCCAGCCGGCTGTG 1518 \vdash CAGCCAGCCTGCTCTG 16 0 Score 12.8; Pred. No. 2 Mismatches 8; DB 1; 2.5e+02; ches 2; Length 17; Indels 0 Gaps 0

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RESULT 351
ABN06184
ABN06184 standard; DNA; 17
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Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. 26-MAY-2000; 2000US-207456P. 21-SEP-2000; 2000US-234687P. 25-MAY-2001; Homo sapiens. Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6176. 29-MAY-2002 WO200192524-A2 2001WO-US16981. (first entry)

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RESULT 352
ABN06185
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04-OCT-2000;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                 of hgDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hgDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hgDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hgDMLP proteins, as specific blomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hgDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hgDMLP-1 may be used for diagnosing a disorder associated with the expression of hgDMLP-1, in particular heart and skeletal muscle disorders hgDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hgDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                   N.B. The sequence data for this patent did not specification, but was obtained in electronic at ftp.wipo.int/pub/published_pct_sequence.
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Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart;
                              Human GDMLP-1
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                                                                                               ABN06185
                                                                                                                           ABN06185 standard; DNA; 17
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Pred. No. 2.5e+02;
0; Mismatches 2;
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ACAAAGACAAGCCCAA 1799 ACAGAGCCAAGCCCAA 16

Matches

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30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
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                                                                                                                      specification, but
                                                                                                                                            N.B. The sequence data
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27-SEP-2000;
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substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and. for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders, hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification enhanced the production acids in samples, as amplification
screening invention.
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30-JAN-2001;
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30-JAN-2001;
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27-SEP-2000;
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2000GB-0024263.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 therapy; vaccine; heart disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Chen W,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shannon ME
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ABN06321
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30-JAN-2001;
30-JAN-2001;
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      protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering
                                                                                                              New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human
                                                            The present invention describes a human genome-derived myosin-like
                                                                                 Disclosure;
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27-SEP-2000;
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skeletal muscle
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14; Conserv
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                                                                                SEQ ID 6313; 214pp;
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2001WO-US00667.
2001WO-US00668.
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Pred. No. 2
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                                                                                English
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therapy; vaccine; heart di
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                                                                                                                                                                              Shannon
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variants

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RESULT 355
ABN06539
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Best Local
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21-SEP-2000;
27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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2000GB-0024263.
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                                                            Penn SG,
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Pred. No. 2.5e+02;
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                                                            Rank DR,
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and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in
                                                                      N.B. The sequence data specification, but was
                                                                                                                                         particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises myosin-like protein hGDMLP-1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein 1 (himmle-1). The protein and polynuclectide sequence hGDMLP-1 can be used in gene therapy and vaccine production. hGDMLP-1 nucleic acids can be used as probes to detect, chara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure;
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                                                                   for this patent did not obtained in electronic
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В Ś Matches Query Match Best Local 1359 GCCTGGAAGAGAAAAG 1374 14; N Similarity GCCTGGTAGAGACAAG Conservative 0.9%; 17 0, Score 12.8; I Pred. No. 2.5e 0; Mismatches 2.5e+02; ches 2; DB 1; Length 17; Indels 0, Gaps 0

Sequence 17

ftp.wipo.int/pub/published_pct_sequence.

BP; 6 A; 3 C; 6 G; 2 T;

0 other;

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RESULT 356
ABN06540;
                    ABN06540 standard; DNA;
                    17
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29-MAY-2002

(first entry)

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:6532.

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

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30-JAN-2001;
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     30-JAN-2001;
30-JAN-2001;
                       30-JAN-2001;
30-JAN-2001;
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                                          27-SEP-2000;
                                                21-SEP-2000;
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                                                                                                       Homo sapiens.
     2001WO-US00663.
2001WO-US00664.
2001WO-US00665.
                       2001WO-US00661.
2001WO-US00662.
                                   2000US-236359P.
2000GB-0024263.
                                               2000US-207456P.
2000US-234687P.
                                                                  2001WO-US16981.
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New polypeptide, for raising antibodies that recognize hGDMLP-1

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RESULT 357
ABNO7353
ID ABNO73
XX ABNO73
AC ABNO73
XX 29-MAY
XX 29-MAY
XX Human;
XX Hum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 14
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30-JAN-2001;
30-JAN-2001;
05-FEB-2001;
                                                                                                                                                                     Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                        Human GDMLP-1
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2001WO-US00669.
2001WO-US00670.
2001US-266860P.
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Pred. No. 2.5e+02;
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06-DEC-2001 WO200192524-A2

ABN07354

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                                                                                                                           The present invention describes a human genome-derived myosin-like CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of CC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 nucleic acids can be used as probes to detect, characterise CC hGDMLP-1 nucleic acids can be used as probes to detect, characterise CC and quantify hGDMLP-1 nucleic acids in samples, as amplification CC substrates, to provide initial substrates for the recombinant engineering CC of hGDMLP-1 protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may CD be used as immunogens to raise antibodies that specifically recognise CC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP proteins, as specific CC disjunction, as therapeutic supplement in patients having specific CC deficiency in hGDMLP-1 production, and in vaccines or for replacement CC diagnosing a disorder associated with the expression of hGDMLP-1, in CC particular heart and skeletal muscle disorders. hGDMLP-1 may be used for CC chromosome 22. The present sequence represents an oligomer used in the greening of the hGDMLP-1 sequence in the exemplification of the present
                                                             Matches
                                                                           Query Match
Best Local
                                                                                                                                                    specification, but was obtained in electron
at ftp.wipo.int/pub/published_pct_sequence.
                                                                                                                                                                     N.B. The sequence data for this patent did not specification, but was obtained in electronic f
                                                                                                                           Sequence 17
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30-JAN-2001;
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27-SEP-2000;
                                                                                                                                                                                                      invention.
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 AAACAGTTGGAAGAAG
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                                                             Conservative
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Pred. No. 2.
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Human GDMLP-1 17-mer scanning 29-MAY-2002 (first entry)

SEQ ID

NO:5

sequence

SEQ

IJ

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of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart, and skeletal muscle disorders hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
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21-SEP-2000;
27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering
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surface-enhanced laser desorption ionization, compri
myosin-like protein hGDMLP-1 -
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                                                                                         nvention.
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Best Local Similarity

0.9%; 87.5%;

Score 12.8; DB 1; Pred. No. 2.5e+02;

Length 17;

Query Match

Sequence

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                        The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantity hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption deficiency in hGDMLP-1 production, and in vaccines or for replacement the harmonic proteins and for replacement and for repl
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therapy.
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The polynucleotide sequences encoding
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2001WO-US00662.
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       hGDMLP-1 may be used for
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of
                                                           New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1 -
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Pred. No. 2.5e+02;
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Best Local S
Matches 14
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27-SEP-2000;
04-OCT-2000;
30-JAN-2001;
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                                                                                                                                                                                                                                     skeletal muscle
                                                                                                                                                                                                                                                                    Human;
                                                                                                                                                                                                                                                                                                   Human GDMLP-1 17-mer scanning
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                                                                  26-MAY-2000;
                                                                                                   25-MAY-2001; 2001WO-US16981
                                                                                                                                     06-DEC-2001.
                                                                                                                                                                      WO200192524-A2
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                                                                                                                                                                                                                                                                                                                                                                      ABN09084
                                                                                                                                                                                                                                                                                                                                                                                                     ABN09084 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    screening of the hGDMLP-1 sequence in the exemplification of the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ftp.wipo.int/pub/published_pct_sequence.
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                                                                                                                                                                                                   sapiens.
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                                                                                                                                                                                                                                                                  genome-derived myosin-like protein 1; GDMLP-1;
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                                                                                                                                                                                                                                                   myosin;
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                  2000US-207456P.
2000US-234687P.
2000US-236359P.
2000GB-0024263.
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                                                                                                                                                                                                                    chromosome 22; gene therapy; vaccine; heart disease; disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    describes a human genome-derived myosin-like
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Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                   SEQ ID NO:5 sequence SEQ ID NO:9076
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
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t directly from WIPO
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RESULT 363
ABN10708
ID ABN107
XX
ABN107
XX
ABN107
XY
DT 29-MAY
XX
DE Human
XX
DE Human;
XW
Human;
XW
Muscle
XX
Skelet

29-MAY-2002

(first entry)

ABN10708 standard; DNA;

17

0;

muscle;

myosin;

skeletal

Human; genome-derived myosin-like protein 1; GDMLP-1;

myosin; chromosome 22; gene therapy; vaccine; muscle disorder; amplicon; screening; ss.

Human GDMLP-1 17-mer scanning

SEQ ID NO:5

sequence

SEQ

heart disease; hGDMLP-1; heart; ID NO:10700.

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                                                                                                                                                                                                                                 cc and quantify himself in uncleic acids in samples, as amplification of the provide initial substrates for the recombinant engineering cc of hGDMLP-1 protein variants having desired phenotypic improvements, and cf or expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption concentration, as therapeutic supplement in patients having specific concentration, as therapeutic supplement in patients having specific consistion, as therapeutic supplement in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in consistency in holds a decided muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the sorresing of the hGDMLP-1 sequence in the exemplification of the present
                                                                  Matches
                                                                                  Query Match
Best Local
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30-JAN-2001;
30-JAN-2001;
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05-FEB-2001;
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1 -
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                                                                                                                                   Sequence
                                                                                                                                                                                   N.B. The sequence data for this patent did not specification, but was obtained in electronic i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, character
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present
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                                                                                                                                                                                                                      invention.
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                        1398 CCTGGAGATAGCCATT 1413
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                                                                                    Similarity
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CCTGGAGACATCCATT 16
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                                                                                                                                   BP; 4 A;
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                                                                  Conservative
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2001WO-US00663.
2001WO-US00664.
2001WO-US00665.
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2001WO-US00667.
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87.5%;
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                                                                                    Score 12.8;
Pred. No. 2
                                                                    Mismatches
                                                                                                                                     0 other;
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t directly from WIPO
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                                                                  Gaps
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Homo sapiens

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                                                                                                                  CC protein 1 (homeles). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The characterise complete in the sequences of hGDMLP-1 nucleic acids can be used as probes to detect, characterise complete in the control of the protein control of the complete in the recombinant engineering confugnation to rations having desired phenotypic improvements, and confugnation variants having desired phenotypic improvements, and confugnation to raise antibodies that specifically recognise concentration and/or amount specifically of hGDMLP-1 proteins, as specific hGDMLP-1 proteins, as specific concentration, as therapeutic supplement in patients having specific concisation, as therapeutic supplement in patients having specific consistation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for charmosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present
                                                                       Query Match
Best Local
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21-SEP-2000;
27-SEP-2000;
                                                                                                               Sequence
                                                                                                                                                        N.B. The sequence data for this patent did not specification, but was obtained in electronic f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID 10700; 214pp; English
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                                                                                                                                                                                        invention.
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2001WO-US00665.

2001WO-US00666.

2001WO-US00668.

2001WO-US00668.

2001WO-US00669.
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2000US-236359P.
2000GB-0024263.
2001WO-US00661.
2001WO-US00662.
2001WO-US00663.
                                                      Conservative
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                                                                    0.9%;
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                                                                                                             G; 1 T; 0 other;
                                                Pred. No. 2.5c
0; Mismatches
                                                                    Score 12.8; DB Pred. No. 2.5e+(
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                                                                      .5e+02;
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                                                                                                                                                        form part of the printed ormat directly from WIPO
                                                                                  Length 17;
                                                      Indels
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                                                     Gaps
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RESULT 364

at ftp.wipo.int/pub/published_pct_sequence

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CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of CC hGDMLP-1 can be used in gene therapy and vaccine production. The CC hGDMLP-1 nucleic acids can be used as probes to detect, characterise CC and quantify hGDMLP-1 nucleic acids in samples, as amplification CC substrates, to provide initial substrates for the recombinant engineering CC of hGDMLP-1 protein variants having desired phenotypic improvements, and CC for expressing the proteins. The hGDMLP-1 proteins or polypeptides may CC hGDMLP-1 proteins, as standards in assays used to determine the CC concentration and/or amount specifically of hGDMLP, proteins, as specific biomolecule capture probes for surface-enhanced laser desorption CC ionisation, as therapeutic supplement in patients having specific CC deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for CC diagnosing a disorder associated with the expression of hGDMLP-1, in CC particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the greening of the hGDMLP-1 sequence in the exemplification of the present
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 invention.

N.B. The sequence data for this patent did not consistent on, but was obtained in electronic consistent.
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                                                                                                                                                                                                                                                                                                                                           The present invention describes a human genome-derived myosin-like
                                                                                                                                                                                                                                                                                                                                                                           Disclosure; .SEQ ID
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30-JAN-2001;
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30-JAN-2001;
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2001WO-US00670.
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RESULT 365
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The invention relates to a nucleic acid molecule (I) which down regulates CC expression of an Ets-related gene (ERG). (I) is useful for treating CC conditions selected from cancer, lymphoma, Ewing's sercoma, melanoma, CC neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca CC neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca CC vulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge CC weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-rendu CC syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for CC treating a patient having a condition associated with the level of ERG, CC by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies CC under conditions suitable for the treatment. Leukaemia or tumour CC angiogenesis is treated by administering (I) to the patient in CC conjunction with one or more of other therapies such as radiation or CC conjunction with one or more of other therapies such as radiation or CC conjunction with one or more of other therapies such as radiation or CC cll, by contacting the cell with (I). (I) is useful for cleaving RNA of CC ERG gene, by contacting (I) with RNA, in the presence of a divalent
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Best Local Similarity 87.9
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber
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ABK17887 standard; RNA; 17
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Pred. No. 2.5e+02;
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                                                                                                                                                                               Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber
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Pred. No. 2.5e+02;
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The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retrinopathy, macular degeneration, neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca yulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge

down regulates

Claim 4; Page 81; 149pp; English.

Weber syndrome, Kippel-Trenaunay-Weber syndrome, Oslusyndrome, leukaemia, osteoporosis and wound healing.

Osler-Weber-rendu ing. (I) is useful

for ERG

condition associated

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the treatment. The method comprises the use of one or more therapies cunder conditions suitable for the treatment. Leukaemia or tumour cangiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting the cell with (I). (I) is useful for cleaving RNA of CR gene, by contacting (I) with RNA, in the presence of a divalent cation such as Mg2+. (I) is useful for diagnosis of conditions and classes related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically cargeting genes that share homology with ERG gene or ERG fusion genes. CRAN17354-ABK2719 represent nucleic acids, including antisense and cenzymatic nucleic acid molecules which regulate expression of ERG, and carried PCR primers of the invention.
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                                                                                       Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Web
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                                                                                                                                                                                                       Von
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cells of the patient with (I) under conditions suitable
                                                                                                                                                                                                   Carlowitz I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        syndrome, leukaemia; osteoporosis; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%;
                                                                                                                                                                                                   McSwiggen JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytostatic; antitumour; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G; 3 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. 2.56
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 12.8; DB Pred. No. 2.5e+
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.5e+02;
2;
                                                                                                                                                                                                   Mclaughlin F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>,</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                         Sturge Weber
                                                                                                                                                                                                   Randi AM
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The invention relates to a nucleic acid molecule (I) which down regulates

The invention relates to a typing kit for judging human leukocyte antigen

Claim 4; Page 90; 149pp; English.

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RESULT 368
ABL31521/c
ID ABL315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC treating a patient having a condition associated with the level of ERG, CC by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies CC under conditions suitable for the treatment. Leukaemia or tumour CC angiogenesis is treated by administering (I) to the patient in CC conjunction with one or more of other therapies such as radiation or CC chemotherapy treatment. (I) is useful for reducing ERG activity in a CC cell, by contacting the cell with (I). (I) is useful for cleaving RNA of CR gene, by contacting (I) with RNA, in the presence of adivalent CC cation such as Mg2+. (I) is useful for diagnosis of conditions and CC classes related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically CC targeting genes that share homology with ERG gene or ERG fusion genes. CABK17354-ABK22719 represent nucleic acids, including antisense and CC enzymatic nucleic acid molecules which regulate expression of ERG, and conjunction and conjunction and conjunction and conjunction and conjunction and conjunction and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.9%;
Best Local Similarity 87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                  Human leukocyte antigen (HLA) typing, useful for of individuals e.g. by determining immunogenetic transplanting between them -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 conditions selected from cancer, lymphoma, Ewing's sarce tumour angiogenesis, diabetic retinopathy, macular degen neovascular glaucoma, myopic degeneration, arthritis, ps vulgaris, angiofibroma of tuberous sclerosis, port-wine weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber syndrome, Osler-Weber syndrome, Cosler-Weber syndrome, C
                                                               Claim 10; Page 284; 345pp; Japanese.
                                                                                                                                                                                                                                                                                   WPI; 2002-122074/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUN-2000; 2000JP-0164798
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUN-2001; 2001WO-JP04662.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; human leukocyte antigen; HLA; genotype; polymorphism;
immunogenetic; transplantation; genetic disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human HLA genotyping oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL31521;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                            (NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2094 CTACCAGCTGGCCAGA 2109
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                                                                                                                                                                                                                                                                                                                                                      Kagiya T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                         Ichihara
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                      Matsumura
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macular degeneration,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NO 1010
                                                                                                                                                                                                                                                                                                                                                      Moriya
                                                                                                                                                                           judging HLA genotypes differences when
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Osler-Weber-rendu
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                                                                                                                                                                                                                                                                                                                                                      Nishida
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Query Match
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given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acide, polypeptides, vectors containing the nucleic acide, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (ABL30512-ABL31809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals.
                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense; sense; tumour; cell degeneration; cancer; Alzheimer schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; antisense; sense; tumour; cell degeneration; cancer; Alzheimer
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                                                                                                                                                                                                                                                         The invention relates to a novel isolated 17 mer nucleic acid sequence,
                                                                                                                                                                                                                                                                                                     Disclosure; Page 40; 720pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-313353/30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MOLE-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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Pred. No. 2.
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les 2;
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                            given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence, with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell
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Pred. No. 2.5e+02;
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cancer but also Alzheimer's

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RESULT 371
ABT34848/c
ID ABT34848 standard; DNA; 17
The invention relates to a novel isolated 17 mer nucleic acid sequence, CG given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel the complement of any of them, or the corresponding RNA. The novel generated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of agency chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these
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2.5e+02;
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RESULT 372
ABT35043/c
The invention relates to a novel isolated 17 mer nucleic acid sequence, consecutive nucleotides from the 17 mer sequence, a sequence with, after coptimal alignment, at least 80 % identity to the 17 mer sequence, a coptimal alignment, at least 80 % identity to the 17 mer sequence, a coptimal alignment, at least 80 % identity to the 17 mer sequence, a coptimal alignment at least 80 % identity to the 17 mer sequence, a coptimal alignment of any of them under highly stringent conditions, or the complement of any of them under highly stringent conditions, or the complement of any of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, coption of recombinate polypeptides. Any of the nucleic acid, coption and for production of recombinant polypeptides. Any of the nucleic acids, copypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral copyperides that are characterised by development of tumours or cell copyperides and specifically cancer but also Alzheimer's disease and schizobrenia. Analysis of the expression of the 17 mer nucleic acids in copypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein
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human fukutin;
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Pred. No. 2
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2.5e+02;
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RESULT 373
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            CC given in the specification, a sequence containing at least 15
CC consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel consolated mucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, cc e.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, cc polypeptides, vectors containing the nucleic acids, cc polypeptides, vectors containing the nucleic acids, cells containing the cc vector or antibodies directed against the polypeptides are useful for cc preparation of pharmaceuticals for prevention and/or treatment of viral cdiseases that are characterised by development of tumours or cell cdiseases that are characterised by development of tumours or cell cdiseases that are characterised by development of tumours or cell containing the sequence of the 17 mer nucleic acids in containing the polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein characterised by development of these containing the invention can be used in gene containing the containing the polypeptides can also be used to generate antibodies, and contains of protein contains o
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therapy. This
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human fukutin; ds.
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Pred. No. 2.5e+0
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         CC given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after coptimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or cc sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers corresponding RNA is probes and primers of the component of a gene chip, in vitro as (anti) sense reagents, can for production of recombinant polypeptides. Any of the nucleic acids, collypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral cliescases that are characterised by development of themours or cell cliescases that are characterised by development of themours or cell capatient samples is useful for diagnosis and/or prognosis of these conficients are useful as components of protein chips. The polypeptides can also be used to generate antibodies, and confice the polypeptide and antibodies are useful as components of protein charge. This polymucleotide sequences of the invention can be used in gene therapy. This polymucleotide sequence represents a tumour suppression
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RESULT 375
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Sequence 17

BP; 6

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4 G;

4 T; 0 other

Best Local Similarity

0.9**%;** 87.5**%**;

Score 12.8; Pred. No. 2

Length 17;

Query Match

17

4.

A 4 C;

w ç

6 T;

0 other; 8; DB 1; 2.5e+02;

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RESULT 376
ABT35577
ID ABT355
XX ABT355
XX Cytost
XX Antise
SX Homo s
XX Hom
                                                       The invention relates to a novel isolated 17 mer nucleic acid sequence, CC given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after coptimal alignment, at least 80 % identity to the 17 mer sequence, a cc sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel cisolated nucleic acids of the invention are useful as probes and primers condetecting, identifying, quantifying and/or amplifying a nucleic acid, cc and for production of recombinant polypeptides. Any of the nucleic acids, ccl as one component of a gene chip, in vitro as (anti)sense reagents, cc and for production of recombinant polypeptides. Any of the nucleic acids, colypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral cliesases that are characterised by development of tumours or cell capeneration, specifically cancer but also Alzheimer's disease and conjugation to separate antibodies is useful for diagnosis and/or prognosis of these conjugations are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene conjugated human fukutin oligonucleotide of the invention.
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Best Local Similarity

Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 175; 720pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated nucleic acid,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polypeptides, antibodies and transfected cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-SEP-2001; 2001FR-0011978.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-SEP-2002; 2002WO-IB04208.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tumour suppression related human fukutin oligo SEQ ID No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-JUN-2003 (first entry
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TTGTTCTACAGCAGAT 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cleic acid, useful for treating viral diseases tumors and cell degeneration, also related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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Pred. No. 2
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. 2.5e+02;
2;
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RESULT 377
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                Query Match
Best Local (
   Matches
                                                                                                                                                                                                                                              given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell
                                                                                                                                        degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for diagnosts and/or prognosts of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABT36389 standard; DNA; 17
                                                                     Sequence 17
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              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention relates to a novel isolated 17 mer nucleic acid sequence,
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                  Similarity
                                                                                                        human
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                                                                                                        This polynucleotide sequence represents a tumour numan fukutin oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 269; 720pp; French.
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   Conservative
                                                                       BP; 5
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                0.9%;
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                                                                       G; 6 T; 0 other;
Score 12.8; DB 1;
Pred. No. 2.5e+02;
0; Mismatches 2;
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                                 Length 17;
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2525 AGCAGTTGGTAGAAGA 2540

Matches Query Match

Local

Similarity

0.9%;

Conservative

<u>,,</u>

Score 12.8; DB 1; Pred. No. 2.5e+02; 0; Mismatches 2;

8; ... 2.5e+02; 2;

Length 17; Indels

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                                     CC given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after CC optimal alignment, at least 80 % identity to the 17 mer sequence, a containing at least 15 coptimal alignment, at least 80 % identity to the 17 mer sequence, a containing at least 15 coptimal alignment, at least 80 % identity to the 17 mer sequence, a coptimal alignment, at least 80 % identity to the 17 mer sequence, a coptimal alignment of any of them under highly stringent conditions, or the complement of any of the mention are useful as probes and primers for detecting, identitying, quantifying and/or amplifying a nucleic acid, copy and for production of recombinant polypeptides. Any of the nucleic acids, collypeptides, vectors containing the nucleic acids, rector or antibodies directed against the polypeptides are useful for copparation of pharmaceuticals for prevention and/or treatment of viral conseases that are characterised by development of tumours or cell containing the expression of the 17 mer nucleic acids in containing the expression of the 17 mer nucleic acids in containing the expression of the 17 mer nucleic acids in containing the expression of the sequences of these containing the collection and containing the collection and containing the expression of the 17 mer nucleic acids in components of these collections and containing the collection of the polypeptides can also be used to generate antibodies, and containing the collection of the polypeptide and antibodies are useful as components of these collections. The nucleic acid sequences of the invention can be used in gene containing the collection of the invention can be used in gene containing the collection of the invention can be used in gene containing the invention can be used i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells -
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Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a novel isolated 17 mer nucleic acid sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 398;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Telerman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-SEP-2001; 2001FR-0011978.
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BP; 7
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A.
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1 C; 5 G; 4 T; 0 other;
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ATCAGTTGGTAAAAGA 17

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RESULT 379
ABT37821/c
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                                                                                                                                                                CC given in the specification, a sequence containing at least 15
CC consecutive nucleotides from the 17 mer sequence, a sequence with, after coptimal alignment, at least 80 % identity to the 17 mer sequence, a cc sequence that hybridizes to them under highly stringent conditions, or CC the complement of any of them, or the corresponding RNA. The novel cisolated nucleic acids of the invention are useful as probes and primers CC for detecting, identifying, quantifying and/or amplifying a nucleic acid, ce.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acid, cc golypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for c proparation of pharmaceuticals for prevention and/or treatment of viral cdiseases that are characterised by development of tumours or cell containing the schizophrenia. Analysis of the expression of the 17 mer nucleic acids in CC diseases. The polypeptides can also be used to generate antibodies, and coths the polypeptide and antibodies are useful as components of these caid sequences of the invention can be used in gene cherapy. This polynucleotide sequence represents a tumour suppression crelated human fukutin oligonucleotide of the invention.
                                                                    Matches
                                                                                   Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                                                        Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel isolated 17 mer nucleic acid sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 438; 720pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Telerman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-SEP-2001; 2001FR-0011978.
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                                 1695 GGAGTTTCCAAGAGAT 1710
 17
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                                                                                       Similarity
                                                                                                                                      BP; 6 A; 4 C; 2 G; 5 T; 0 other;
                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amson R,
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                                                                                  0.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tuijnder M;
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                                                                                  Score 12.8; DB 1;
Pred. No. 2.5e+02;
                                                                 Mismatches
                                                                                                  Length 17
                                                                   Indels
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RESULT 380
ABT3800AC
ID ABT380
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ABT380
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ABT380
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ABT380
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ABT380
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                                                                                                                                                                                                                                                                                                                                                                            CC given in the specification, a sequence containing at least 15
CC consecutive nucleotides from the 17 mer sequence, a sequence with, after coptimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel complement of any of them, or the corresponding RNA. The novel conditions is contained by the invention are useful as probes and primers corresponding RNA to a nucleic acid, or detecting, identifying, quantifying and/or amplifying a nucleic acid, or and for production of recombinant polypeptides. Any of the nucleic acid, and for production of recombinant polypeptides and containing the vector or antibodies directed against the polypeptides are useful for diseases that are characterised by development of tumours or cell containing the nucleic acid, and preparation, specifically cancer but also Alzheimer's disease and component of the sequence of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein characterise acid sequences of the invention can be used in generate antibodies. The nucleic acid sequences of the invention can be used in generate antibodies.
                                                                                                                           Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                              Sequence 17 BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABT38000 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                            related human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel isolated 17 mer nucleic acid sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 459;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated nucleic acid, useful for treating viral disea associated with tumors and cell degeneration, also related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-313353/30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MOLE-) MOLECULAR ENGINES LAB.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            schizophrenia;
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                                                          2533 GTAGAAGACTTGGATC 2548
16
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                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                        polynucleotide sequence represents a tumour suppression fukutin oligonucleotide of the invention.
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                                                                                                                                                       0.9%;
                                                                                                                                                                                                                                                          7 C; 1 G; 5 T; 0 other;
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                                                                                                                           ٥,
                                                                                                                                                          Score 12.8;
Pred. No. 2.
                                                                                                                                  Mismatches
                                                                                                                                                              .5e+02;
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                                                                                                                               Indels
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RESULT 381

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                                                                         Query Match
Best Local S
Matches 14
                                                                                                                                                                                      degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                       given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence, with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by dependent of tumours or conditional degeneration. Specifically cancer but also Alzhaimer's disease and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells
                                                                                                                                                      Sequence 17 BP; 5
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                                                                   Local Sim-
                                    1377
                                                                                           Similarity
                                  GATTACAGCTTCCCCA 1392
GATCACAGCTGCCCCA 16
                                                                           Conservative
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                                                                                                                                                      A; 7 C; 3 G; 2 T; 0 other;
                                                                                           0.9%;
87.5%;
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                                                                                             Score 12.8; DB
Pred. No. 2.5e+(
                                                                           Pred. No. 2.5
); Mismatches
                                                                                           8; DB 1;
2.5e+02;
                                                                                                               Length 17;
                                                                           Indels
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RESULT 382 ABT38046 ID ABT380

ABT38046 standard; DNA; 17

RESULT 383 ABT39470/c ID ABT394 XX AC ABT394

ABT39470 standard; DNA; 17

ВP

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ABT39470

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CC consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel complement of any of them, or the corresponding RNA. The novel complement of any of the invention are useful as probes and primers of retecting, identifying, quantifying and/or amplifying a nucleic acid, ce g, as one component of a gene chip, in vitro as (anti)sense reagents, can for production of recombinant polypeptides. Any of the nucleic acids, collas containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for comparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell containing the separation, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in CR patient samples is useful for diagnosis and/or prognosis of these containing the polypeptides can also be used to generate antibodies, and the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene containing the contained human fukutin oligonucleotide of the invention.
                                                                        Matches
                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells -
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                                                                                                                                                  Sequence 17 BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel isolated 17 mer nucleic acid sequence,
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                                   2318 ATCAGAGTGATGTCTG 2333
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                                                                      14;
N
 ATCAGAGTGAAGTATG 17
                                                                        Conservative
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                                                                                                                                                  Α,
                                                                                        0.9%;
                                                                                                                                                  1 C; 6 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tuijnder M;
                                                                        0
                                                                                         Score 12.8; DB 1;
Pred. No. 2.5e+02;
                                                                          Mismatches
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                                                                                                           Length 17;
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RESULT 384
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                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel isolated 17 mer nucleic acid sequence, CC given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or cc the complement of any of them, or the corresponding RNA. The novel cisolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, ce.g. as one component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, cells containing the nucleic acids, cells containing the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of themours or cell caparistion, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in cellicates. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of these chips. The nucleic acid sequences of the invention can be used in gene therapy. This polyrucleotide sequence represents a tumour suppression can be used in gene therapy. This polyrucleotide sequence represents a tumour suppression can be used in gene therapy. This polyrucleotide sequence represents a tumour suppression can be used in gene therapy. This polyrucleotide sequence represents a tumour suppression can be used in gene therapy. This polyrucleotide of the invention can be used in gene therapy. This polyrucleotide of the invention can be used in gene therapy. This polyrucleotide of the invention can be used in gene therapy. This polyrucleotide of the invention can be used in gene therapy. This polyrucleotide of the invention can be used in gene therapy.
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Best Local
12-JUN-2003
                                            ABT39771;
                                                                                   ABT39771 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                A; 1 C; 4 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                            0.9%;
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                                                                                                                                                                                                                                                                                                               Score 12.8; DB 1;
Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                   2.5e+02;
2;
                                                                                                                                                                                                                                                                                                                                 Length 17;
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Tumour suppression related human fukutin oligo SEQ ID No
                                    5408
   gene
chip;
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Čýtostatic; virucide; neuroprotective; nootropic; neuroleptic; antisense; sense; tumour; cell degeneration; cancer; Alzheimer' schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds. œ disease;

Homo sapiens.

WO2003025175-A2

27-MAR-2003

17-SEP-2002; 2002WO-IB04208

17-SEP-2001; 2001FR-0011978

(MOLE-) MOLECULAR ENGINES LAB

Telerman Þ Amson R, Tuijnder M;

WPI; 2003-313353/30

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells -

Disclosure; Page 666; 720pp; French.

CC given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after coptimal alignment, at least 80 % identity to the 17 mer sequence, a coptimal alignment of any of them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel consisted nucleic acids of the invention are useful as probes and primers copy of them, or the corresponding RNA. The novel copy of the invention are useful as probes and primers copy as one component of a gene chip, in vitro as (anti)sense reagents, copy as one component of a gene chip, in vitro as (anti)sense reagents, copy and for production of recombinant polypeptides. Any of the nucleic acids, copy peptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for corparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tunours or cell diseases that are characterised by development of tunours or cell containing the 17 mer nucleic acids in copy of the sequence of the 17 mer nucleic acids in copy of the polypeptides can also be used to generate antibodies, and copy peptides are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene contained the copy of the components of protein copy of the copy of t The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15

Sequence 17 BP; 2 A; 5 C; 2 G; 8 T; 0 other;

Query Match Best Local S Matches 14 14; Similarity Conservative 0.9%; 0, Score 12.8; DB 1; Pred. No. 2.5e+02; o. Mismatches 2; Length 17; Indels 0 Gaps

0

S 1364 GAAGAGAAAAGGAGAT 1379

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17

GCAGAGAAAATGAGAT 2

RESULT 385 ABT39947/c ID ABT399

ABT39947 standard; DNA; 17

13-JUN-2003 (first entry)

Tumour suppression related human fukutin oligo SEQ ID No

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RESULT 386
ACA06845/c
SX SX PX PX PX B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence with after comptimal alignment, at least 80 % identity to the 17 mer sequence, a containing at least 15 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, ce.g. as one component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, ce.g. as one component of a gene chip, in vitro as (anti)sense reagents, ce golypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tunours or cell caparation, specifically cancer but also Alzheimer's disease and components of these containing the sequences of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these containing the polypeptides can also be used to generate antibodies, and coips. The nucleic acid sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene containing the collectide sequence represents a tumour suppression containing the containing the collectide of the invention can be used in gene containing the containing the containing the collectide of the invention can be used in gene containing the containing the containing the collectide of the invention can be used in gene containing the cont
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associated with tumors and cell degeneration, also related
polypeptides, antibodies and transfected cells
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llarity 87.5%;
Conservative
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0; Mismatches
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Pred. No. 2.5e+02;
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ches 2;
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Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;

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1398 CCTGGAGATAGCCATT 1413

Query Match Best Local S Matches 14

Similarity

0.9%; ilarity 87.5%; Conservative

0;

Mismatches

Score 12.8; DB 1; Pred. No. 2.5e+02;

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Length 17; Indels

0,

Gaps

0

14;

Sequence 17

ВP;

3 A;

6 C; 4 G; 4 U; 0 other;

NFKB sub-unit modulating inozyme substrate

#664

03-JUN-2003

(first entry)

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CC (1) is useful for cleaving RNA comprising a sequence of REL-A gene, in Ct the presence of a divalent cation, especially Mg^2-+. The enzymatic and cantisense nucleic acid molecules are useful for treating breast, lung, CC prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cc cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or cmultidrug resistant cancer. The method involves use of other drug complete such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cc gemeitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as the chematoid arthritis, restenosis, asthma, Crohn's disease, diabetes, cobesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-AUG-1994;
07-DEC-1992;
18-MAY-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inczyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 3; Page 36; 72pp; English.
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96US-0777916.
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regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inozyme, Zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and cantisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, ossophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug characters such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, generitable or radiation therapy. The enzymatic and antisense nucleic conditions are such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, generitable or radiation therapy. The enzymatic and antisense nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; osophageal cancer; stomach cancer; bladder cancer; parceatic cancer; corvical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; sementabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; Page 40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases
                                                                                                                                                                                                                                                                                                                                                         The invention describes an enzymatic nucleic acid molecule (I) which down
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-340953/32
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18-MAY-1994;
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92US-0987132.
94US-0245466.
96US-0777916.
radiation therapy. The are also useful for tre
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                       The invention relates to a novel stabilising reaction reagent for use in the amplification and/or detection of a target nucleic acid comprising: preparing a reaction mixture with e.g. a nucleic acid as template, at least 1 primer and RNaseH; and incubation of the reaction mixture for a defined period of time to form a reaction product during the amplification of such target nucleic acid. The method is useful for stabilising and long-term storage of reaction reagents for highly sensitive and specific amplification and detection of nucleic acids particularly in identifying pathogenic microorganisms or viruses in a sample using chimeric oligonucleotide primers, which is useful in genetic engineering and clinical medicine. This polynucleotide sequence represents an oligo relating to the novel stabilising reaction reagent method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes obesity, autoimmune disease, lupus, multiple sclerosis, transplant, rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel
                                                                                                                                                                                                                                                                                       Method for stabilizing and storing reaction reagents for specific amplification and detection of nucleic acids particularly in e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stabilising reaction reagent; PCR; primer; RNaseH; long-term specific amplification; pathogenic microorganism; chimeric; genetic engineering; clinical medicine; ss.
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20-AUG-2001; 2001JP-0249689.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Stabilising reagent method related oligo SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                Uemori T,
                                                                                                                                                                                                                                                                          pathogenic microorganisms
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ilarity 87.5%;
Conservative
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Cato I;
                                                                                                                                                                                                                                         177pp;
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                                                                                                                                                                                                                                            Japanese
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Pred. No. 2.
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                                                                                                                                                                                                                                                                            or viruses
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2.5e+02;
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Best Local
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06-JUN-2001;
10-SEP-2001;
                                                                                                                                                  acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-F human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-Heumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ6216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66530 - ABZ66585 represent substrate/target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                          Claim 58; Page 97; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                   Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encodi HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
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                        2151 TTTAGCAGCCAGAAAT 2166
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                                                                  Similarity
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UUUAGUAACCAGAAAU 17
                                                                                                                                        for the human ribozymes of the invention.
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; 2001US-296249P.
; 2001US-318471P.
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Pred. No. 2.5e+02;
0; Mismatches 2;
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Pred. No. 2.5e+02;
4; Mismatches 2;
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RESULT 390 ABZ60832

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Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
                                                                                                                       ABZ61775;
                                                                                                                                                   ABZ61775
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
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06-JUN-2001;
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                                                            Human H-Ras DNAzyme target #566.
                                                                                           21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 8 A; 2 C; 5 G; 2 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                             sequences for the human ribozymes of the invention
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                                                                                                                                                                                                                                                          2190 GATGAAAATAGCAGAC 2205
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                                                                                                                                                                                                                                                                                         13;
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                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                   standard; RNA; 17
                                                                                                                                                                                                                              GAAGAAUAUAGCAGAC 17
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; 2001US-296249P.
; 2001US-318471P.
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                                                                                                                                                                                                                                                                                                     Score 12.8; DB 1;
Pred. No. 2.5e+02;
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                                                                                                                                                                                                                                                                                                                     DB 1;
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RESULT 392
ABZ62111/c
ID ABZ62111;
XX
AC ABZ62111;
XX
DT 21-MAR-2003 (first entry)
XX
DE Human H-Ras DNAzyme target #9
XX
DE Human, ribozyme; short interf
XX
Human, ribozyme; cancer; AIDS;
XX
Human trheumatic; cancer; AIDS;
XX
PD 05-DEC-2002.
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PD 05-DEC-2002.
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PP 29-MAY-2001; 2001US-294140P.
PR 29-MAY-2001; 2001US-296249P.
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Best Local Similarity 68.8%;
Matches 11; Conservative
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                                                                                                                                                                                            Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
       29-MAY-2001; 2001US-294140P.
06-JUN-2001; 2001US-296249P.
                                                                                                                                                                                                                                                                Human H-Ras DNAzyme target #902
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
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06-JUN-2001; 2001US-296249P
10-SEP-2001; 2001US-318471P
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Pred. No. 2.5e+0
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2.5e+02;
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06-JUN-2001; 2001US-296249P
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                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC
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Pred. No. 2.
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Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences

WPI; 2003-140484/13.

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RESULT 394
ABZ65478/C
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Best Local (
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                                                  The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Rahuman immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and
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10-SEP-2001; 2001US-318471P.
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anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung
                                                                                                                                                        Claim 4; Page 151; 185pp; English.
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Best Loc Matches Query Match

Local

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0.9%; llarity 92.3%; Conservative

Score 12.6; I Pred. No. 1.9e 1; Mismatches

1.9e+02; DB 1; 0

Length 13;

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Gaps

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RESULT 395
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                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.

ABC00010-ABC99989, ABF00010-ABF99893, ABH00010-ABH99989 and ABI00010-ABH829898, The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     prostate, bladder, or pancreatic cancer, and HIV infection, and AID: The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention.
    Sequence
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                                         ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID 41027; 29pp + Sequence Listing; German
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llarity 87.5%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
    3 A;
  1 C;
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Pred. No. 2.5e+02;
0; Mismatches 2;
    5 T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              7,
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      1 other;
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TTGGATCGAATTY

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RESULT 396
ABC41011/c
ID ABC410
XX ABC410
XX ABC410
XX ABC410
XX SNP; s
XW Peptid
XW SNP; s
XW Peptid
XW Centra
XX WO2001
XX WO1; 2
XX WPI; 2
XX This i
CC Canda cy
CC Coligon
CC ABC000
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ABC7
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                                                                                                                                                                                                                                                       RESULT 397
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                               76782/c
ABC76782 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.

ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI32073 represent the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                 Oligonucleotide SEQ ID NO 76799 for detecting SNP TSC0019618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                21-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID 41028; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    methylation status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-657177/75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      hes 12;
                                                                                                                                                                                                                                                                                                                                                                                 2542 TTGGATCGAATTC 2554
                                                                                                                                                                                                                                                                                                                                           13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                             (first entry)
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Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
.9e+02;
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RESULT 398
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.

ABC00010-ABC99989, ABP00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fig. wipo.int/pub/published_pct_sequences.
                                                                                            WO200177384-A2
                                                                                                                           Homo sapiens.
                                                                                                                                                           central nervous
                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
                                                                                                                                                                                                                        Oligonucleotide SEQ ID NO 76800 for detecting SNP TSC0019618
                                                                                                                                                                                                                                                         21-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13 BP; 3 A; 0 C; 2 G; 7 T; 1 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    methylation status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                        ABC76783 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-APR-2000; 2000DE-1019173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                           system;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.9%;
                                                                                                                                                         gastrointestinal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Berlin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 12.6; DB 1;
Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ea. No. 1.9e+02;
Mismatches 0
                                                                                                                                                           respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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07-APR-2000; 2000DE-1019173 06-APR-2001; 2001WO-IB00713.

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RESULT 399
ABC79816/c
ID ABC798
XX ABC798
XX ABC798
XX Oligon
XX SNP; s
KW peptid
KW centra
XX Homo s
XX H
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                               Claim 1; SEQ ID 79833;
                                                                                            methylation status
                                                                                                                      Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single nucleotide polymorphisms and cytosin
                                                                                                                                                                                                                     WPI; 2001-657177/75
                                                                                                                                                                                                                                                                                                                                                                                                     07-APR-2000; 2000DE-1019173
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12; Conserv
                                                                                                                                                                                                                                                                                                                                            EPIGENOMICS
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                                                                                                                                                                                                                                                                               Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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92.3%;
                               29pp + Sequence Listing; German
                                                                                                                                                                                                                                                                                  Berlin K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 12.6; DB Pred. No. 1.9e+(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6; DB 1;
1.9e+02;
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RESULT 400
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ABC00010-ABC99989, ABF0010-ABF9989, ABH0010-ABH9989 and ABI00010-ABH82073 represent the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell designed to detect single nucleotide polymorphisms and methylation status -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide SEQ ID NO 79834 for detecting SNP TSC0020266.
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Pred. No. 1.
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABC00010-ABC99989, ABF00010-ABF99989, ABC00010-ABC99989, ABF00010-ABF99989, ABC00010-ABC99989, ABC00010-ABC9989, ABC00010-ABC9989, ABC00010-ABC9989, ABC00010-ABC9989, ABC00010-ABC9989, ABC00010-ABC9989, ABC00010-ABC9989, ABC00010-ABC9989, ABC00010-ABC9989, ABC00010-ABC99989, ABC00010-ABC99989, ABC00010-ABC99989, AB

ftp.wipo.int/pub/published_pct_sequences.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a

Claim 1; SEQ ID 79834; 29pp + Sequence Listing; German

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RESULT 401
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                                                                                                                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.

ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99999 and ABH00010-ABH99989, The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
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                                                                                                        Sequence 13
                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID 184727; 29pp + Sequence Listing; German.
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12; Conserv
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                                                 0.9%;
ilarity 92.3%;
Conservative
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                                                            Score 12.6; DB 1;
Pred. No. 1.9e+02;
                                                    Mismatches
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RESULT 402 ABF84731/c

Homo sapiens

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RESULT 403
ABH29758/c
ID 9ABH297
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AC ABH297
XX
AC ABH297
XX
Oligon
XX
Oligon
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SNP; S
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ABC00010-ABC99989, ABF00010-ABF99889, ABH00010-ABH9989 and ABI00010-ABG99899, are present the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                Oligonucleotide SEQ ID NO 229735 for detecting
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les 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A; 6 C; 0 G; 2 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                      clarity 92.3%;
Conservative
                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                      1;
                                                                                                                                                                                                                                                                                                                                                                      Score 12.6; DB 1;
Pred. No. 1.9e+02;
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RESULT 404
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ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention:
NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
          Olek A,
                                                            07-APR-2000;
                                                                                    06-APR-2001;
                                                                                                                18-OCT-2001
                                                                                                                                                                   Homo
                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
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          Berlin K;
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Pred. No. 1.9e+02;
1; Mismatches 0
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RESULT 405
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ABC00010-ABC99999, ABF00010-ABF99989, ABF00010-ABF99989 and ABF00010-ABF99989, ABF00010-ABF99989 and peresent the oligomers described in the invention.
NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosiae methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a

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                                                                                                          range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.

ABC00010-ABC9989, ABF00010-ABF9989, ABF00010-ABH99989 and ABI00010-ABH393073 represent the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
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                                                                 Sequence 13
                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID 261822; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                  methylation status
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide SEQ ID NO 261822 for detecting SNP TSC0063523
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0.9%;
ilarity 92.3%;
Conservative
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Score 12.6; DB 1;
Pred. No. 1.9e+02;
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RESULT 407
ABK55502/c
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specific condition or disease predicted to be associated with SELL activity. The method is also useful in screening for compounds targeting SELL to treat a specific condition or disease predicted to be associated with SELL activity, e.g. detecting which of the SELL haplotypes or haplotype pairs present in individual members of a population with the specific disease of interest enables one to screen for compounds that display the highest desired agonist or antagonist activity for each of the most frequent SELL isoforms present in the disease population. A polymorphic variant of SELL is useful in studying the effect of the variation on the biological activity of SELL, on the binding affinity of candidate drugs targeting SELL for the treatment of neonatal pertussis (whooping cough) and in assays to measure the binding affinities of one or more candidate drugs targeting the SELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel genetic variants of selectin L lymphocyte adhesion molecule 1 (SELL) gene useful for therapeutic purposes and for expressing SELL protein useful in identifying drugs to treat whooping cough -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligonucleotide; ss.
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RESULT 408
AAD26040/c
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Query Match
Best Local Similarity
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Best Local
                                                                               familial dysbetalipoproteinaemia, type III hyperlipoproteinaemia, atherosclerosis, and Alzheimer's disease. They are useful to validate APOE as a candidate agent for treating a specific condition or disease predicted to be associated with APOE activity and in the design of clinical trials of candidate drugs for treating a specific condition or disease predicted to be associated with APOE activity. Genotyping or haplotyping methods are useful to screen for compounds targeting or haplotyping methods are useful to screen for compounds targeting APOE to treat a specific condition or disease associated with APOE activity. The present DNA sequence is an allele specific oligonucleotide (ASO) probe which is used for detecting human APOE gene polymorphisms.
                                                                                                                                                                                                                                       The patent discloses novel genetic variants of human apolipoprotein E (APDE) gene. The invention also relates to compositions and methods for haplotyping and/or genotyping the APDE gene. The haplotyping methods of the invention are useful for improving the efficacy and reliability of several steps in the discovery and development of drugs for treating diseases associated with APDE activity, e.g.
                                                                                                                                                                                                                                                                                                                                                                                      Genotyping human apolipoprotein gene of individual for determining haplotype of individual, involves determining identity of nucleotide pair at specific polymorphic sites for two copies of gene -
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; antilipaemic; neuroprotective; nootropic; genetic variant; APOE; apolipoprotein E; haplotyping; familial dysbetalipoproteinaemia; therapy; genotyping; type III hyperlipoproteinaemia; Alzheimer's disease; atherosclerosis; polymorphism; allele specific oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein. ABK55465-ABK55559 represent SELL gene allele-specific
oligonucleotides of the invention.
                                                                                                                                                                                                                                                                                                                                                          Claim 16; Page 14; 78pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Choi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-APR-2000; 2000US-197188P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-APR-2001; 2001WO-US12303
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les 12; Conserv
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Pred. No. 2.3e
1; Mismatches
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      Pred.
                     Score
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                                                      T;
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    12.6; DB 1;
No. 2.3e+02;
                                                      other;
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2.3e+02;
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RESULT 410
AAV95609
ID AAV956
XX
AC AAV956
XX
AC AAV956
DT 24-FEB
XX
DE Human
XX

AAV95609

standard;

RNA;

14

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24-FEB-1999

(first

entry)

Human c-fos target sequence nucleotide position

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RESULT 409
AAQ41008
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                               Query Match
Best Local Similarity
Matches 13; Conserv
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                                                                                                                                       Modified eukaryotic cytoplasmic DNA virus prodn. - involdirect molecular cloning of modified DNA molecule contgcytoplasmic DNA virus genome
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                                                                   Sequence 14 BP; 3 A;
                                                                                            The synthetic early/late promoter selP was used to express gp160-gene of the HIV-1 MN strain. The final plasmid is
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                                                                                   pselP-gp160MN
                                                                                                                      Example; Fig 9.5.B; 206pp; English
                                                                                                                                                                                   WPI; 1993-126461/16.
                                                                                                                                                                                                    Dorner F,
                                                                                                                                                                                                                                      26-AUG-1991;
20-JUL-1992;
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                                                                                                                                                                                                                                                                                                  AU9221269-A
                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                     Plasmid; cloning;
                                                                                                                                                                                                                                                                                                                                                                                               Sequence around translational start codon
                                                                                                                                                                          P-PSDB; AAR34645.
                                                                                                                                                                                                                     (IMMO ) IMMUNO AG.
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\mathbf{\mu}
                CCGTGGCCGTGAAG 1824
                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
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CCATGGCCGTGAAG 14
                                                                                                                                                                                                     Falkner FG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                 Conservative
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92US-0914738.
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                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                       restriction site;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                        0.9%;
                                                                   4 C; 5 G; 2 T; 0 other;
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                                         Score 12.4; DB 1;
Pred. No. 2.2e+02;
                                  Mismatches
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                                                 Length
                                   Indels
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AAV97202/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes an enzymatic nucleic acid molecule which specifically cleaves RNA derived from a c-fos gene. AAV95401 to AAV95540 and AAV95541 to AAV95584 represent hammerhead ribozymes and hairpin ribozymes, respectively, which specifically cleave human c-fos. AAV95261 to AAV95400 and AAV95585 to AAV95628 represent human c-fos target sequences. The enzymatic nucleic acid molecules can be used for treating cancer associated with elevated levels of c-fos oncogene, especially leukaemias, neuroblastomas and lung, breast and colon cancers. The ribozymes may also be used as diagnostic tools to examine genetic drift and mutations within diseased cells, or to detect the presence of c-fos RNA in a cell.
24-NOV-1997;
28-JAN-1997;
                                       14-JAN-1998;
                                                                  30-JUL-1998
                                                                                            WO9832843-A2
                                                                                                                        Solanum tuberosum.
                                                                                                                                                 flower formation;
                                                                                                                                                            Solanidine; glucosyltransferase; potato; citrate synthase; target; hammerhead ribozyme; hairpin ribozyme; alkaloid biosynthesis;
                                                                                                                                                                                                     Potato citrate synthase target sequence position 539.
                                                                                                                                                                                                                                  01-MAR-1999
                                                                                                                                                                                                                                                              AAV97202;
                                                                                                                                                                                                                                                                                      202/c
AAV97202 standard; RNA; 14 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic nucleic acid molecules which derived from a c-fos gene - useful for to levels of c-fos, especially cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-JAN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mutation; diseased cell; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; c-fos; hammerhead ribozyme; hairpin ribozyme; target site; cancer; oncogene; leukaemia; neuroblastoma; diagnosis; genetic drift;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9832846-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        McSwiggen JA,
                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 5 A; 3 C; 4 G; 2 U; 0 other;
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97US-0979416.
97US-0036545.
                                       98WO-US00738
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                                                                                                                                                 cleavage; solanaceous plant; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                          78.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                             Score 12.4; DB Pred. No. 2.2e+
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                                                                                                                                                                                                                                                                                                                                                                                                                          2.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    specifically cleave RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      treating conditions related
                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 14;
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CC alkaloids; or (ii) involved in flower formation. AAV95322 to AAV96334, CC and AAV96335 to AAV96354 represent potato solanidine glucosyltransferase CC hammerhead and hairpin ribozymes, respectively. AAV95325 to AAV959981, CC and AAV96335 to AAV96773 represent potato solanidine glucosyltransferase CC target sequences. AAV96773 to AAV97170, and AAV97171 to AAV97195 (C respectively. AAV96773 to AAV97170, and AAV97171 to AAV97195 (C respectively. AAV96773 to AAV97172, and AAV97196 to AAV97220 represent CC potato citrate synthase target sequences. Ribozymes of the present CC invention can be used to inhibit the synthesis of toxic alkaloids in CC solanaceous plants, particularly potato but also tomato, pepper, CC aubergine and ditura or to inhibit flowering in potato, lettuce, spinach, CC aubergine and turf grass. Also the ribozymes can be used for RNA CC manipulation in the same way that restriction endonucleases are for DNA, CC sweet potato and turf grass. Also the ribozymes can be used for RNA CC manipulation in the same way that restriction endonucleases are for DNA, CC as well as to examine genetic drift and mutations in plants and to consensus sequences within a family of related genes, and being CC cataluric manipulation and to a respect at a family of related genes, and being consensus sequences within a family of related genes, and being consensus sequences within a family of related genes.
Sequence 14 BP; 3 A; 5 C; 1 G; 5 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes enzymatic nucleic acid molecules with RNA-cleaving activity (e.g. ribozymes) which are capable of modulating the expression of plant genes: (i) involved in biosynthesis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New enzymatic nucleic acid(s) - useful biosynthesis or regulating flowering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-427939/36.
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                                                  very low
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Best Local S
Matches 13
                                                                                                                       Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiinflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome;
                                                                                                                                                                                                                                                        Human TIE-2 target site SEQ ID NO:2432.
                                                                                                                                                                                                                                                                                    19-JUN-2000
                                                                                                                                                                                                                                                                                                                 AAA19206;
                                                                                                                                                                                                                                                                                                                                           AAA19206 standard; RNA; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                              2473 ATGATGATGAGGGA 2486
                                                                                                                                                                                                                                                                                                                                                                                                                 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                 ATGATGATCAGGGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.9%;
                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 12.4; DB 1;
Pred. No. 2.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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0,

24-MAR-1999; 07-OCT-1999 WO9950403-A2

99WO-US06507

Homo sapiens

Kippel-Trenaunay-Weber

syndrome; Osler-Weber-Rendu

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RESULT 413
AAA09609
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA23422 represent their corresponding target sequences. The ribozymes of the invention are used for modulating the synthesis, expression and/or stability of an mRNA encoding angiogenic factor, especially ARWT, integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARMD), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2, integrin subunit alpha-6, or integrin subunit beta-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA1675 to AAA1767 and AAA1763 to AAA17622 represent ribozyme sequences for RNT, and AAA17168 to AAA17630 and AAA17623 to AAA1764 represent their corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086 and AAA19155 to AAA20361 and AAA21501 to AAA21505 represent tribozyme sequences; AAA19223 to AAA20361 and AAA21501 to AAA21505 to AAA21500 and AAA21505 to AAA21689 to AAA21688 represent their corresponding target sequences; AAA21506 to AAA21688 represent their corresponding target sequences; AAA21506 to AAA21688 represent their corresponding target sequences; AAA21506 to AAA22475 and AAA23263 to AAA23342 represent ribozyme sequence for integrin subunit beta 3, and AAA22476 to AAA2362, AAA23343 to
                  24-MAR-2000; 2000WO-US07979
                                                                                               WO200056901-A2
                                                                                                                                       Synthetic
                                                                                                                                                                               primer;
                                                                                                                                                                                             Linear expression element; circular expression element; LEE; CEE;
gene function; antigen; promoter function; vaccine; antibody production;
                                                                                                                                                                                                                                                            Primer SEQ ID 3 used in LEE and CEE production.
                                                                                                                                                                                                                                                                                                    29-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                 AAA09609 standard; RNA; 14 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 14 BP; 4 A; 1 C; 4 G; 5 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 56; Page 139; 305pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel ribozymes for modulating the synthesis, expression and/or stability of an mRNA encoding an angiogenic factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2198 TAGCAGACTTTGGA 2211
                                                                                                                                                                               88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   UAGCAGAUUUUGGA 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 12.4; DB 1;
Pred. No. 2.2e+02;
4; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Coeshott C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        McSwiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
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RESULT 414
AAA09611/c
ID AAA096
XX
AC AAA096
XX
DT 29-JAN
XX
DE Primer
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression element (LEE or CEE). The method comprises obtaining a DNA segment consisting of an open reading frame (ORF) and linking the ORF to a promoter and a terminator. The LEEs and CEEs are useful for screening gene function, biological effects of gene functions, antigens and promoter functions. LEEs and CEEs are useful for producing antibodies to an ORF and also in developing immunological reagents, and may be used in the production of vaccines. Sequences AAA09607-A09611 represent primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Producing a linear or circular expression element useful for a number of molecular biology protocols, comprises obtaining a DNA segment consisting of an open reading frame and linking it to a promoter and terminator
                      Producing a linear or circular expression element useful for a number of molecular biology protocols, comprises obtaining a DNA segment consisting of an open reading frame and linking it to a promoter and
                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                            Linear expression element; circular expression element; LEE;
gene function; antigen; promoter function; vaccine; antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA09611;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA09611 standard; RNA; 14 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-MAR-1999;
31-MAR-1999;
                                                                                  WPI; 2000-628266/60.
                                                                                                           Sykes KF, Johnston
                                                                                                                                                                                   24-MAR-1999;
                                                                                                                                                                                                               24-MAR-2000; 2000WO-US07979
                                                                                                                                                                                                                                           28-SEP-2000
                                                                                                                                                                                                                                                                        WO200056901-A2.
                                                                                                                                                                                                                                                                                                                               primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                      Primer SEQ ID 5 used in LEE and CEE production.
                                                                                                                                                                                                                                                                                                                                                                                                                   29-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disclosed in the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to the production of a linear or circular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 116; 117pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-628266/60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sykes KF,
                                                                                                                                          (TEXA ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     госат
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1882 ATGATGAAGATGAT 1895
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14 BP; 5 A; 0 C; 4 G; 5 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AUGAUGAUGAUGAU 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Johnston SA;
                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                          TEXAS SYSTEM
                                                                                                                                                                     99US-0125864.
99US-0127222.
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99US-0127222.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%;
                                                                                                              SA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 12.4; DB 1;
Pred. No. 2.2e+02;
4; Mismatches 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                             CEE;
                                                                                                                                                                                                                                                                                                                                              production;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Disclosure;

Page 117;

117pp; English.

invention relates to the production of a linear or circular

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RESULT 415
AAA89885
ID AAA898
XX AAA898
XX AAA898
XX DNA se
XX Vaccin
KW Vaccin
KW Gag-po
XX Human
OS Synthe
XX US6103
XX I9-DEC
PR 26-AUC
PR 26-A
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-DEC-1994;
26-AUG-1991;
20-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression element (LEE or CEE). The method comprises obtaining a DNA segment consisting of an open reading frame (ORF) and linking the ORF to a promoter and a terminator. The LEEs and CEEs are useful for screening gene function, biological effects of gene functions, antigens and promoter functions. LEEs and CEEs are useful for producing antibodies to an ORF and also in developing immunological reagents, and may be used in the production of vaccines. Sequences AAA09607-A09611 represent primers
                                                                                                                                          fowlpox virus for generating or priming an immune response against heterologous protein in a vertebrate. Suitable proteins include HIV proteins such as HIV gp160, HIV Gag and HIV Gag-Pol proteins. The proteins sequence was associated with the generation or use of the
                                                                                                                                                                                                                                                                                                                                                                 Use of modified vaccinia virus and priming an immune response against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA89885
                                                                           Sequence
                                                                                                                                                                                                                                             The present invention relates to the use of modified vaccinia virus
                                                                                                                                                                                                                                                                                               Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pfleiderer M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-MAY-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Immunodeficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gag-pol; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA89885 standard; DNA; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disclosed in the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (IMMO) IMMUNO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000-557665/51.
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                                                                                                                                                                                                                                                                                          9; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB15068
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                                                                                                                           viruses
                                                                           14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 5
                                                                           BP; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         around start codon of modified gp160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
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91US-0750080.
92US-0914738.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AG.
                                                                                                                                                                                                                                                                                             99; 171pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Falkner
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     0.9%;
92.9%;
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                                                                           G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scheiflinger F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
     Score 12.4; DB 1;
Pred. No. 2.2e+02;
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Pred. No. 2.2e+02;
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o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              response;
                                                                                                                                                                                                                                                                                                                                                                 fowlpox virus for generating or
HIV gp160, HIV Gag and HIV Gag-Pol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dorner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gp-160;
                           Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 14;
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RESULT 416
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                                                         Matches
                                                                                      Query Match
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                                                                                                                                                                             The invention relates to a method for producing a modified eukaryotic cytoplasmic DNA virus by direct molecular cloning of a modified DNA molecule comprising a modified cytoplasmic DNA virus genome such as a vaccinia virus, containing a heterologous insert encoding a protein. The method involves molecularly cloning the DNA directly into a host cell via a unique restriction endonuclease cleavage site, to be packaged into infectious virions and then recovering them. The host cell is infected with a helper virus for this purpose. The method is useful for producing recombinant proteins and modified eukaryotic cytoplasmic DNA viruses such as poxviruses and iridoviruses found in vertebrates and insects. This
                                                                                                                                                                                                                                                                                                                                                                                                 Producing recombinant protein using modified vaccinia viral expression system, comprises directly cloning heterologous insert encoding protein into the viral genome into unique restriction endonuclease cleavage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          unique restriction endonuclease cleavage site; infectious virion; ds; helper virus; poxvirus; iridovirus; vertebrate; multiple cloning site; human immunodeficiency virus glycoprotein 160MN; gp160MN; mutant.
                                                                                                                     Sequence 14 BP; 3 A; 4 C; 5 G; 2 T; 0 other;
                                                                                                                                                  sequence represents a DNA encoding a modified HIV gp160 peptide used the construction of chimeric vaccinia viruses.
                                                                                                                                                                                                                                                                                                                                                     Example 9; Fig 9.5B; 172pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-AUG-1991;
20-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human Immunodeficiency Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytoplasmic DNA virus; direct molecular cloning; vaccinia virus; insect;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modified gp160MN peptide sequence DNA in vselP-gp160-virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAS12903 standard; DNA; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                               P-PSDB; AAU07642
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dorner F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BAXT ) BAXTER AG.
                                                        Local Similarity nes 13; Conserv
                           1811 CCGTGGCCGTGAAG 1824
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1811 CCGTGGCCGTGAAG 1824
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2001-535006/59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scheiflinger
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                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         91US-0750080
92US-0914738
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                                                                      92.9%;
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                                                                        Score 12.4;
Pred. No. 2.
                                                           Mismatches
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                                                                         2.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 418
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Best Local S
Matches 13
                                                               myocardial ischaemia; Kawasaki disease; myocardial ischaemia; Kawasaki acquired i
                                                                                                                                        gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumen necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; Philadelphia chromosome; inflammation; autoimmune disease;
                   Respiratory Syncytial Virus
                                                                                                         atherosclerosis; myocardial infarction; strotransplant rejection; rheumatoid arthritis;
                                                                                                                                                                                                            Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1;
                                                                                                                                                                                                                                                                 RSV 1B hammerhead ribozyme target sequence (nt. position 242).
                                                                                                                                                                                                                                                                                                    25-MAR-2003
04-APR-1997
                                                                                                                                                                                                                                                                                                                                                        AAT56846;
                                                                                                                                                                                                                                                                                                                                                                                         AAT56846 standard; RNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 14 BP; 3 A; 1 C; 6 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     adenylate cyclase associated primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a light-controlled adenylate cyclase which polypeptide extracted from the duplicate flagellum body of Euglena o partial peptide. The adenylate cyclase and the DNA are used in an in light switch. The current sequence represents a light-controlled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New light-controlled adenylate cyclase, useful for in vivo light switch
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (WATA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-JUL-2000; 2000JP-0240426
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Light-controlled adenylate cyclase; flagellum; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Light-controlled adenylate
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WATANABE M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gracilis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGTTAATGTGGGAG 2358
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0; Mismatches
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Pred. No. 2.2e+02;
                                                                      immune deficiency syndrome;
                                                                                       septic shock; HIV;
                                                                                                                         stroke; restenosis;
                                                                                                         psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 14;
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   RESULT 419
AAX66621/c
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04-APR-1994;
07-APR-1994;
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Modak A,
                                                                                                                                                                                                                Sequence 15
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23-SEP-1994
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15-AUG-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Thompson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-AUG-1994
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17-AUG-1994;
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                                                                                                           2237
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                                                                                                                                                                                                                  BP; 8
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The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves mRNA coding for a protein of respiratory syncytial virus (RSV) at the nucleotide base position indicated in the DE line. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozymes cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes cleave the target sequences and can be used for treatment and diagnosis of RSV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 2; Page 265; 407pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ribozymes having modified bases and methods
for use in inhibiting disease related genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1995-351090/45.
                                                                                      l Similarity
                                        ACTATTACAAAAAG 2250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Karpeisky A, Kisich K,
Pavco P, Beigleman L,
JD, Tracz D, Usman N,
                                                                                        Conservative
                                                                                                                                                                                                                     25-MAR-2003
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94US-0314397.
94US-0316771.
94US-0319492.
94US-0321993.
94US-0334847.
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94US-0357577.
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94US-0293520.
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94US-0291433.
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94US-0271280.
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94US-0222795.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chowrira B, Direnzo A, Draper KG, Dudycz LW;
isky A, Kisich K, Matulic-adamic J, Mcswiggen
P, Beigleman L, Sullivan SM, Sweedler D;
cacz D, Usman N, Wincott FE, Woolf T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PHARM INC
                                                                                                                                                                            P
                                                                                                          71.4%;
                                                                                                                                                                          3 C; 1 G; 3 U; 0 other;
                                                                                                                                 0.9%;
15
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                                                                                                                                                                                                                       correct PI field.)
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•
                                                                                                          Score 12.4; DB 1;
Pred. No. 2.4e+02;
                                                                                      Mismatches
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                                                                                                                               Length 15;
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AAX66621 standard;

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Query Match
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       Matches
                                                                                                                                                               The present invention describes a novel enzymatic nucleic acid (ENA) having a hammerhead motif (HM) comprising: (1) at least 5 ribose residues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) at least ten 2'-C-methyl modifications; and (iv) a 3'-end modification. The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for enhancing graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene expression which accomment treatment with variance and alloantic producing the non-specific effects upon gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-MAY-1995;
04-MAY-1995;
07-JUL-1995;
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13-DEC-1994;
23-DEC-1994;
23-DEC-1994;
17-FEB-1995;
                                                                                                          expression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly specific. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Draper K, Gustofi
Beigelman L, Karı
Matulic-Adamic J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo
                                                                                                                                                                                                                                                                                                                                                                                          Claim 10;
                                                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic nucleic acid molecules having a hammer-head motif - used for the treatment of arthritis, induction of graft tolerance or treatment of auto-immune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-JUL-1995;
07-AUG-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human CD40 hammerhead ribozyme target
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM
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                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gustofson J,
                                                                 BP; 2
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C J, Jarvis T,
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95US-0390850.

95US-0426124.

95US-0432874.

95US-0434509.

95US-0000974.

95US-0512861.
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A, Modak A, Usman N, Bur
s T, Thompson JD, Wincott
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                            Score 12.4; DB
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2.4e+02;
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ncott F;
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RESULT 420
AAT49651/c
ID AAT496
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AC AAT496
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CC AAT49608-T49863 represent target sequences for the human cholesterol
CC ester transfer protein (CETP) hammerhead (HH) ribozymes (see
CA AT498881-T50137). CETP is a 74 kD glycoprotein that facilitates neutral
CC lipid transfer between plasma lipoproteins. The numbering of the targets
CC refers to the position of the cleavage site in full length CETP. The
CC ribozyme binds to 5 nucleotides either side of this site, provided the
CC expression of the mRNA. By inhibiting CETP, thereby blocking synthesis and/or
CC expression of the mRNA. By inhibiting CETP, the reverse cholesterol
CC transport (RCT) pathway can be inhibited (or eliminated) thereby
CC preventing the reduction in size density of the high density lipoproteins
CC (HDL), prolonging HDL half life and therefore increasing HDL levels.
CC The ribozymes can be used to treat conditions associated with abnormal
CC etherosclerosis, peripheral vascular disease, hyperbetalipoproteinaemia,
CC atherosclerosis, peripheral vascular disease, hyperbetalipoproteins of
CC diabetes, transplant, atherectomy and angioplastic restenosis. By
CC inhibiting CETP, the levels of HDL and low density lipoproteins (LDL),
CC and the HDL:LDL ratio are favourably altered (a decrease in LDL levels,
CC and a corresponding increase in HDL levels). The HH ribozymes can also
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New ribozyme(s) for cleaving cholesterol ester transfer protein mRNA - useful for preventing or treating initial development, progression or regression of vascular diseases, esp. familial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     neutral lipid tränsfer; plasma lipoprotein; atherosclerosis; atherectomy; reverse cholesterol transport; high density lipoprotein; therapy; CETP; familial hypercholesterolaemia; dyslipidaemia; hyporalphalipoproteinaemia; peripheral vascular disease; hyperbetalipoproteinaemia; RCT; inhibitor; angioplastic restenosis; low density lipoprotein; diabetes; HDL; human;
                                be used diagnostically to study genetic drift and mutations in diseased cells, and to detect CETP mRNA. As the HH ribozymes target specific regions of the CETP gene, they have low non-specific activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hammerhead ribozyme; cholesterol ester transfer protein; mRNA cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human CETP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-FEB-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Couture L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                      Matches
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Best Local
                                                                                                                                                                                                                                                                                                                The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VBGP). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples
                                                                                                                                                                                                                                 Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 9; Page 191; 218pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Escobedo J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-JAN-1996;
26-OCT-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human flt-1 and KDR hammerhead ribozyme target site #57.
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                                                                                                                                                                                                                                                                                              nucleic acid
                                                      1822 AAGATGTTGAAAGA 1835
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   AAAAUGUUGAAAGA 14
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95US-0005974.
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                                                                                                                                                                                                                                                                                           molecules from the present invention.
                                                                                                                                                                                                                                 A; 0 C; 3 G; 3 U; 0 other;
                                                                                                                                     0.9%;
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                                                                                                                Score 12.4; DB
Pred. No. 2.4e+(
3; Mismatches
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); Mismatches
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Pred. No. 2.4e+02;
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                                                                                                             DB 1;
1.4e+02;
les 1;
                                                                                                                                                                    Length 15;
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RESULT 422 AAT99048/c

RESULT 423 AAZ07078 ID AAZ070

AAZ07078 standard; DNA; 15

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TTTCTGATCTGGTG 1869

Matches

13;

Conservative

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Query Match Best Local Similarity

0.9%;

Score 12.4; DB 1; Pred. No. 2.4e+02; Mismatches

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Length 15;

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Gaps

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This sequence represents a probe for a wild type HIV reverse CC transcriptase (RT) gene fragment. This sequence can be used in the method CC of the invention for determining the susceptibility to antiviral drugs of CC viruses which contain RT genes and are present in a biological sample. It CC comprises: (1) releasing, isolating or concentrating the polynucleic CC acids present with at least one suitable primer pair; (3) hybridising the golynucleic acids of step (1) or (2) with at least two RT gene probes. CC the probes being applied to known locations on a solid support, and are CC capable of simultaneously hybridising to their respective target regions under appropriate hybridisation and wash condition allowing the detection of homologous targets, or with the probes hybridising specifically with a sequence complementary to any of the target sequences; (4) detecting the CC hybrids formed in step (3); and (4) inferring the nucleotide sequence at the codons of interest (codons 38-44, 47-53, 65-72, 73-77, 148-154, CC 180-187, 212-216, and 217-220), and/or the amino acids of the codons of codons of interest and/or antiviral drug resistance spectrum, and possible the type obtained in step (4). The method is specifically used to detect antiviral crug resistant strains of viruses containing RT genes, especially HIV creating the method can also be used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Reverse transcriptase gene; HIV; RT gene; antiviral drug susceptibility; virus susceptibility; antiviral drug resistant viral strain; retrovirus; Hepadnaviridae; HIV RT genotyping; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Determining susceptibility to antiviral drugs of reverse transcriptase containing viruses - useful for genotyping
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 13; Page 39; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-JUN-1996;
26-JAN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immunodeficiency virus type
 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antiviral resistant HIV
                                      HIV RT.
 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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96EP-0870005.
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 5 C; 1 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   induced HIV RT gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genotyping HIV RT
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RESULT 424
AAX57566
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a branched composition (I) which is useful for improving the solubility of synthetic polymers (II) or aids in minimizing or eliminating self-aggregation of (II), where (II) is a nucleic acid (or analogue), peptide, peptide nucleic acid (PNA), polymerde, chimera or a linked polymer. Modification of (II) by (I) can facilitate synthesis, purification and analysis of many insoluble polymers, and particularly purine-rich PNA polymers labeled with hydrophobic labels. The products can be used in research, diagnostic and therapeutic applications. The present sequence represents a PNA and in the account of the present sequence represents a PNA committee of the present sequence represents 
                     AAX57566
                                                           AAX57566 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Branched compositions for improving the solubility of synthetic polymers or minimizing or eliminating polymer self-aggregation, particularly in peptide nucleic acids
                                                                                                                                                                                                                                                                                                                                                              Sequence 15 BP; 5 A; 0 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 12; Page 40; 81pp; English.
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27-JAN-1998;
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                                                                                                                                                                                                                          1882 ATGATGAAGATGAT 1895
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98US-0072772.
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92.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'g is modified to g-E-NH2, uncharged ether modifying
                                                                ВP
                                                                                                                                                                                                                                                                    Pred. No. 2.4(); Mismatches
                                                                                                                                                                                                                                                                                            Score 12.4; DB 1;
Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                         present invention.
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AAZ59271/c
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Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequences AAX57562-X57571 represent antisense oligonucleotides targeted to a region spanning 4-9 codons downstream of the AUG translation initiation codon of the human insulin-like growth factor: 1 receptor (IGF-1R) gene. The antisense oligonucleotides inhibit the expression of IGF-1R, which in turn induces apoptosis, especially in a tumour cell. The oligonucleotides can be administered via a liposome.
                               24-JUN-1998;
19-OCT-1998;
                                                                  23-JUN-1999;
                                                                                        29-DEC-1999.
                                                                                                                                                                    Haemopoietin receptor family; NR8; antibody; diagnosis;
                                                                                                                                                                                                                                                             AAZ59271 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15 BP; 4 A; 4 C; 4 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 16; 23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        oligonucleotides
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         (CHUG-) CHUGAI RES INST MOLECULAR MEDICINE INC
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                                                                                                                                                                                                                                                                                                                     GGAGCCAGACTTCA 14
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inhibition; induction; apoptosis; tumour; liposome;
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                               98JP-0214720.
98JP-0297409.
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Pred. No. 2.4e+02;
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                       The invention relates to the isolation of sequences encoding human haemopoietin receptor protein family NR8 genes. The NR8 family sequences were initially searched comparison on a nucleic acid database with the nucleic acid probe sequence TGGAGYNNNTGGAGY encoding the amino acid sequence Trp-Ser-Xaa-Trp-Ser. The sequences AAZ559258-Z59300 and AAZ90816-Z90925 represent specific examples of probe
              sequences used in the search.
                                                                                                                 Example 1; Page 38; 176pp; Japanese.
                                                                                                                                            Hemopoietin receptor protein family NR8 formation disorders -
                                                                                                                                                                                                            Nomura H,
                                                                                                                                                                                                                                                              24-JUN-1998;
19-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Human NR8 gene probe #16.
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                                                                                                                                                                                                                                      (CHUG-)
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the diagnosis
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98JP-0297409.
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                                                                                                                                                                                                                                                                                                                                                                                                        family; NR8; antibody;
der; fusion protein; pro
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0; Mismatches
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              Antibodies to
 formation disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                          probe;
            the NR8 family proteins
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RESULT 428 AAZ59282/c ID AAZ592 XX

AAZ59282

standard; DNA;

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RESULT 427
AAZ59278/c
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AAZ59278
                                                                                                  haemopoietin receptor protein family NR8 genes. The NR8 family sequences were initially searched for comparison on a nucleic acid database with the nucleic acid probe sequence TGGAGYUNUNTGAGY encoding the amino acid sequence Trp-Ser-Xaa-Trp-Ser. The sequences ARZ59258-Z59300 and ARZ90816-Z59925 represent specific examples of probe sequences used in the search. Antibodies to the NR8 family proteins are used for the diagnosis of blood formation disorders. Compounds identified as binding to the proteins are used for the treatment of such disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                            Example 1;
                                                                                                                                                                                                                                 Hemopoietin receptor protein formation disorders -
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19-OCT-1998;
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                                                                                                                                                                                       The invention relates to the isolation of sequences encoding
                                                                                                                                                                                                                                                                                                                                                             23-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                         (CHUG-) CHUGAI RES
                                                   Local
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 CTCCATGCACTCCA
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                                                                                                                                                                                                            Page 38; 176pp; Japanese
                                                                                  BP; 3
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                                          Conservative
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98JP-0297409.
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                                         Score 12.4; I
Pred. No. 2.46
0; Mismatches
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Pred. No. 2.
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                                                             Length 15;
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RESULT 429
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Best Local
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                    23-JUN-1999;
                                                                     29-DEC-1999.
                                                                                                                       WO9967290-A1
                                                                                                                                                                        Homo
                                                                                                                                                                                                                                              Haemopoietin
                                                                                                                                                                                                                                                                                          Human NR8 gene
                                                                                                                                                                                                                                                                                                                                                24-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                 AAZ59300,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 39; 176pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hemopoietin receptor protein family NR8 used for diagnosis of blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-116933/10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-JUN-1998;
19-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Haemopoietin red
blood formation
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                                                                                                                                                                   sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTCCATGCACTCCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTCCATGAACTCCA 1594
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 3 A; 1 C; 7 G; 4 T; 0 other;
                                                                                                                                                                                                               receptor family; NR8; antibody; diagnosis; ion disorder; fusion protein; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          receptor family; NR8; antibody; diagnosis; ion disorder; fusion protein; probe; ss.
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                                                                                                                                                                                                                                                                                            probe #43
                       99WO-JP03351
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98JP-0297409
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Pred. No. 2.4e+02;
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RESULT 430
AAZ90836/c
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Best Local Similarity
Matches 13; Conserv
The invention relates to the isolation of sequences haemopoietin receptor protein family NR8 genes. The sequences were initially searched for comparison on
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                                                  Example 1; Page 40; 176pp; Japanese
                                                                              Hemopoietin receptor protein formation disorders -
                                                                                                                     WPI; 2000-116933/10.
                                                                                                                                             Nomura H,
                                                                                                                                                                                                 24-JUN-1998;
19-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                               blood
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   as binding to the proteins are used for the treatment of such disorders.
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                                                                                                                                                                                                                                                                                           WO9967290-A1
                                                                                                                                                                                                                                                                                                                                                            Haemopoietin receptor family; NR8; antibody; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                      Human NR8 gene probe
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19-OCT-1998;
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                                                                                                                                              Maeda
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98JP-0297409.
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98JP-0297409.
                                                                                                                                                                                                                                                                                                                                              disorder; fusion protein;
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                                                                                          family NR8 used for diagnosis of blood
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Pred. No. 2.4e+02;
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                                                                                                                                                                                                                                                                                                                                              probe; ss.
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NR8 famil
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AAZ90837/c
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Matches
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                                                                                                                                                                                                   The invention relates to the isolation of sequences encoding human haemopoietin receptor protein family NR8 genes. The NR8 family sequences were initially searched for comparison on a nucleic acid database with the nucleic acid probe sequence TGGAGYNNNTGGAGY encoding the amino acid sequence Trp-Ser-Xaa-Trp-Ser. The sequences the ARZ59258-Z59300 and ARZ90816-Z90925 represent specific examples of probe sequences used in the search. Antibodies to the NR8 family proteins are used for the diagnosis of blood formation disorders. Compounds identified as binding to the proteins are used for the treatment of such disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI;
                                                                                                                                                                 Sequence 15
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19-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    formation disorders
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                                                                                                     Local
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CTCCATGCACTCCA 1
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                                                                                                                                                                 BP; 3 A; 1 C; 7 G; 4 T; 0 other;
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                                                                                Conservative
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98JP-0297409.
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                                                                              Score 12.4; DB
Pred. No. 2.4e+(
0; Mismatches
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                                                                                                                    Length 15;
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RESULT 432
AAZ90846/c
ID AAZ908
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AAZ90861/c
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                                                                                  Human NR8 gene probe
                       Homo sapiens
                                                            Haemopoietin receptor family; NR8; antibody; diagnosis;
                                                                                                              24-MAY-2000
                                                                                                                                                            AAZ90861
                                                                                                                                                                                                                                                                                                                         Sequence 15 BP; 4 A; 0 C; 6 G; 5 T; 0 other;
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19-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human NR8 gene probe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hemopoietin receptor protein family NR8 used for diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CHUG-) CHUGAI RES INST MOLECULAR MEDICINE INC
                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                1581 CTCCATGAACTCCA 1594
                                                 formation
                                                                                                                                                                                                                          14
                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                              standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Maeda
                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                            (first entry)
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98JP-0297409.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99WO-JP03351
                                              disorder; fusion protein;
                                                                                                                                                                                                                                                                                      0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           #74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP
                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                      Score 12.4; DB 1;
Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                            Mismatches
                                               probe;
                                                                                                                                                                                                                                                                                                Length 15;
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AAZ90870/c
ID AAZ908
XX AAZ908
AC AAZ908
XX Human
XX Haemop
KW Haemop
KW Homo s
XX Homo s
YX HOMO 967
XX HOMO 967
XX 24-JUN
PR 19-OC1
XX 24-JUN
PR 19-OC1
XX WPI; ;
XX WPI; ;
XX Hemopo
PT format
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to the isolation of sequences encoding human haemopoietin receptor protein family NR8 genes. The NR8 family sequences were initially searched for comparison on a nucleic acid database with the nucleic acid probe sequence TGGAGYNNNTGGAGY encoding the amino acid sequence Trp-Ser-Xaa-Trp-Ser. The sequences probe AAZ59258-Z59300 and AAZ90816-Z90925 represent specific examples of probe sequences used in the search. Antibodies to the NR8 family proteins are used for the diagnosis of blood formation disorders. Compounds identified as binding to the proteins are used for the treatment of such disorders.
               Hemopoietin receptor protein family NR8 used for diagnosis of blood formation disorders -
                                                       WPI; 2000-116933/10
                                                                               Nomura H,
                                                                                                                                    24-JUN-1998;
19-OCT-1998;
                                                                                                                                                                            23-JUN-1999;
                                                                                                                                                                                                      29-DEC-1999.
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                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                    Haemopoietin red
                                                                                                                                                                                                                                                                                                                            Human NR8
                                                                                                                                                                                                                                                                                                                                                     24-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                           AAZ90870 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hemopoietin receptor protein family NR8 used for diagnosis of blood formation disorders - {\sf rec}
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19-OCT-1998;
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                                                                                                          (CHUG-) CHUGAI RES INST MOLECULAR MEDICINE
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les 13; Conser
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                                                                                                                                                                                                                                                                                                                          gene
                                                                                 Maeda M;
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ilarity 92.9%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 3
                                                                                                                                                                                                                                                                                receptor family; NR8; antibody; diagnosis; ion disorder; fusion protein; probe; ss.
                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                    98JP-0214720.
98JP-0297409.
                                                                                                                                                                                                                                                                                                                          probe
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98JP-0297409.
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Pred: No. 2.4e+02;
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RESULT 435
AAZ90883/c
ID AAZ908
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                                                                   The invention relates to the isolation of sequences encoding human haemopoietin receptor protein family NR8 genes. The NR8 family sequences were initially searched for comparison on a nucleic acid database with the nucleic acid probe sequence TGGAGYNNNTGGAGY encoding the amino acid sequence Trp-Ser-Xaa-Trp-Ser. The sequences AAZ59258-Z59300 and AAZ90816-Z59925 represent specific examples of probe sequences used in the search. Antibodies to the NR8 family proteins are used for the diagnosis of blood formation disorders. Compounds identified as binding to the proteins are used for the treatment of such disorders.
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                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                      24-JUN-1998;
19-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human NR8 gene probe #111.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 as binding to the proteins are used for the treatment of such disorders.
                                                                                                                                                                                                                                                                Hemopoietin receptor protein family NR8 used
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                                                                                                                                                                                                                                                     formation disorders
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13; Conserv
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nilarity 92.9%;
Conservative
                                           BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     receptor family; NR8; antibody; diagnosis;
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98JP-0297409
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0.9%;
                                           1 C; 7 G; 4 T; 0 other;
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Pred. No. 2.4e
O; Mismatches
Score 12.4; DB 1;
Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein;
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             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       probe;
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             Length
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Matches

Conservative

0

Mismatches

Indels

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Gaps

0

Best Local Similarity

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AAZ90902/
ID AAZ9
XX
AC AAZ9
XX
DT 24-M
XX
DE Huma:
XX
XX
Haem
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AAZ90895/c
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                                                                                                                       RESULT 437
                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                 Query Match
Haemopoietin receptor family; NR8; antibody; diagnosis;
                         Human NR8
                                               24-MAY-2000
                                                                       AAZ90902;
                                                                                              AAZ90902 standard; DNA; 15 BP
                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                98
                                                                                                                                                                                                                                                                                           sequences used in the search. Antibodies to the NR8 family proteins are used for the diagnosis of blood formation disorders. Compounds identified
                                                                                                                                                                                                                                                                                                                sequences were initially searched for comparison on a nucleic acid database with the nucleic acid probe sequence TGAGYNNTGGAGY encoding the amino acid sequence Trp-Ser-Xaa-Trp-Ser. The sequences AAZ59258-Z59300 and AAZ90816-Z90925 represent specific examples of probe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JUN-1998;
19-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Haemopoietin receptor family; NR8; antibody; diagnosis; blood formation disorder; fusion protein; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human NR8 gene probe #123
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                                                                                                                                                                                                                                                                                                                                                                    haemopoietin receptor
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                                                                                                                                                          CTCCATGCACTCCA 1
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                       gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maeda M;
                                                                                                                                                                                                                                                                               to the
                                                                                                                                                                                                                                                         BP; 3 A; 1 C; 7 G; 4 T; 0 other;
                                                                                                                                                                                                         Conservative
                                              (first entry)
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                       probe
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98JP-0297409.
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                                                                                                                                                                                                                                                                              proteins are used for the treatment of such disorders.
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                                                                                                                                                                                                                   0.9%;
                         #130
                                                                                                                                                                                                                                                                                                                                                                  protein family NR8 genes. The NR8 family
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0; Mismatches
                                                                                                                                                                                                                   Score 12.4; DB 1;
Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                         for diagnosis of
                                                                                                                                                                                                                               Length 15;
                                                                                                                                                                                                          Indels
                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                         blood
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RESULT 438
AAZ90906/c
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Best Local Similarity
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                        Nomura H,
                                                                                                              24-JUN-1998;
19-OCT-1998;
                                                                                                                                                                                                                                  29-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                         plood
                                                                                                                                                                                                                                                                                                                                                                                               Haemopoietin
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19-OCT-1998;
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                                                                                                                                                                                    23-JUN-1999;
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                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hemopoietin receptor protein family NR8 used for diagnosis of blood
formation disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nomura H,
                                                                   (CHUG-) CHUGAI RES INST MOLECULAR MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CHUG-) CHUGAI RES INST MOLECULAR MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1581
                                                                                                                                                                                                                                                                                                                                                                         formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14 CTCCAGGAACTCCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTCCATGAACTCCA 1594
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Maeda M;
                        Maeda M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 2 A; 3 C; 6 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                               receptor family; NR8; antibody; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                   98JP-0297409
                                                                                                                                       98JP-0214720
                                                                                                                                                                                    99WO-JP03351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98JP-0214720.
98JP-0297409.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99WO-JP03351
                                                                                                                                                                                                                                                                                                                                                                         disorder; fusion protein; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                               #134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12.4; DB 1;
Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of such
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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WPI;

2000-116933/10

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RESULT 439
AAZ399088 ctandard; DNA; 15 BP.
XX
AC AAZ39908;
XX
AC AAZ399908;
XX
DT 24-MAY-2000 (first entry)
XX
DE Human NR8 gene probe #136.
XX
Haemopoietin receptor family; NR
KW blood formation disorder; fusion
XX
CS Homo sapiens.
XX
PN W09967290-A1.
XX
PN W09967290-A1.
XX
PN W29-DEC-1999; 98W9-JP03351.
XX
PN W29-DEC-1998; 98JP-0214720.
PR 23-JUN-1998; 98JP-0214720.
PR 23-JUN-1998; 98JP-027409.
XX
PN W09967290-A1.
XX
PN 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Example 1; Page 44; 176pp; unpublic to the isolation of sequences encoding numbers has protein family NR8 genes. The NR8 family has sequences were initially searched for comparison on a nucleic acid database with the nucleic acid probe sequence TGGAGYNNNTGGAGY encoding the amino acid sequence TTp-Ser-Xaa-TTp-Ser. The sequences the sequence Ttp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-TTp-Ser-Xaa-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hemopoietin receptor protein family NR8 used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CHUG-) CHUGAI RES INST MOLECULAR MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          haemopoietin receptor protein family NR8 genes. The NR8 family sequences were initially searched for comparison on a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hemopoietin receptor protein family NR8 used formation disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               росат
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disorder; fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%;
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Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NR8; antibody; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.4e+02;
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RESULT 440
AAH26020/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                              07-JUN-1996;
30-SEP-1993;
30-SEP-1994;
The present sequence is that of a stem-loop rat Syk antisense oligonucleotide (ODN). The loop domain consists of 3 tandemly joined antisense ODNs (see AAH26012-14), each targeted to a different region of rat Syk mRNA. Thus, the stem-loop antisense ODN can hybridise to 3 different sites of rat Syk mRNA. It was used in an experiment to demonstrate inhibition of histamine release from histamine-containing RBL-2H3 rat mast cells using Syk antisense ODNs. The invention provides a claimed method of inhibiting the release of a mediator from a Syk-producing cell by introducing into the cell an antisense construct that targets an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 2
                                                                                                                              Example 6; Fig 9; 35pp; English.
                                                                                                                                                   Inhibiting the release of a mediator from a Syk-producing cell, in gene therapy for treating inflammatory conditions or asthma, introducing into the cell Syk antisense oligonucleotides
                                                                                                                                                                                                   WPI; 2001-380484/40.
                                                                                                                                                                                                                        Schreiber AD,
                                                                                                                                                                                                                                                                    07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                   05-JUN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            stem_loop
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  stem_loop
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Syk; tyrosine kinase; rat; antisense; asthma; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Stem-loop
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH26020;
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                                                                                                                                                                                                                                                                                                                            14-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                          US6242427-B1
                                                                                                                                                                                                                                                                                                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antiasthmatic;
                                                                                                                                                                                                                                             AIND (-SAKD)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1581 CTCCATGAACTCCA 1594
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                                                                                                                                                                                                                                                                   96US-0657884.
93US-0129381.
94US-0316425.
95US-0483530.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammation;
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58..60
/*tag=
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36..53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligonucleotide targeting rat Syk
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                                                                                                                                                                                                                                                                                                                                                                                               "phosphorothioate linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                "phosphorothioate linkage'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12.4;
Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phagocytosis;
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                                                                                                                                                                           useful
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RESULT 441
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                                                                The patent relates to polymorphic variants of human cholinergic receptor, comuscarinic 3 (CHRM3) gene. The polymorphic variant comprises at least one single nucleotide polymorphisms selected from cytosine at PS1, adenine at PS2 or PS3, and cytosine at PS4. The invention also relates to a method for genotyping and haplotyping the CHRM3 gene for identification of cyariants. The polymorphic variant is useful for therapeutic purposes, for studying the expression and biological function of CHRM3. CC for studying the expression and biological function of CHRM3 protein.

CC The variant is also useful in diagnostics and forensic applications. CC The recombinant nonhuman organism transfected with the polymorphic variant is useful for studying expression of CHRM3 isogenes in vivo, for in vivo screening and testing of drugs targetted against CHRM3 CC for in vivo screening and testing of drugs targetted against CHRM3 CC for Alzheimer's disease, Sjogren's syndrome, disorders associated with smooth muscle contractility and sudden infant death syndrome. The CHRM3 content is useful in drug screening asserts and its arribodies are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; cholinergic receptor muscarinic 3; CHRM3; drug screening, single nucleotide polymorphism; forensic application; gene therapy; Alzheimer's disease; Sjogren's syndrome; smooth muscle contractility; sudden infant death syndrome; genotyping; haplotyping; ASO; sudden infant death syndrome; genotyping; haplotyping; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Syk encoding sequence such that inhibition is effected. Also claimed is a method of treating an inflammatory condition in a patient by administering an antisense construct that targets Syk encoding sequences and inhibits Syk kinase production.
                                                                                                                                                                                                                                                                                                                                                                           Claim 15; Page 19; 54pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                purposes
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel polymorphic variant of reference sequence for human cholinergic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-300326/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-OCT-2000; 2000WO-US28247
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-JUL-2001
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                     samples.
                                                        protein variant is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-OCT-1999;
                                  useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GENA-)
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primer used for
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               ariant is useful in drug screening assays and its antibodies immunoassays to detect CHRM3 protein variants in biological The present sequence is an allele-specific oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                 muscarinic 3 gene, useful for diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Denton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nandabalan K,
detecting human CHRM3 gene polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      muscarinic 3 gene ASO primer #11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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Pred. No. 2.4e+0
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RESULT 443
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AC AAF695

AAF69549 standard; DNA; 15

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AAF69549

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RESULT 442
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                                                                                                              (Mitochondrial, proton carrier) (UCP3) gene and polymorphisms. The polymorphisms are associated with obesity, especially diabetes mellitus associated obesity. They polymorphism may be identified and analysed to determine whether an individual is susceptible to obesity and may be used as the basis for targeted design of drugs to treat obesity. The present sequence was used the identification and amplification of UCP3 polymorphisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                         Nucleic acids encoding uncoupling protein 3 (mitochondrial, proton carrier) (UCP3) proteins comprising single nucleotide polymorphisms useful for the design of drugs for treating obesity -
                                                                                        Sequence
                                                                                                                                                                                                                    Claim 15; Page 21; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                       08-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                              08-SEP-2000; 2000WO-US24784.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UCP3; uncoupling protein 3; polymorphism; obesity;
diabetes mellitus; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        UCP3 polymorphism
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                                                                                                                                                                                             The present invention relates to the human uncoupling protein
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                     1444 ATGGTGGTAACAGT 1457
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                                                                                                                                                                                                                                                                                                             Choi JY,
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                                                                                          BP; 4 A; 2 C; 5 G; 4 T; 0 other;
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                                            Conservative
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                                                       Score 12.4; DB 1
Pred. No. 2.4e+02
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Pred. No. 2.
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                                             Mismatches
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                                                                  DB 1;
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RESULT 444
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XX JO-MAR
XX JGFBP2
XX ANTISE
KW CYLOST
KW SKIN d
KW IGF MOT
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Best Local S
Matches 13
                                                 Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation, psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; seleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to polymorphisms of the human interleukin 4 receptor-alpha gene (IL4R-alpha; see AAF57718 for the reference sequence). Polymorleotides comprising polymorphic gene variants are useful for therapeutic purposes. For example, where a patient may benefit from expression of a particular IL4Ralpha protein isoform, an expression vector encoding the isoform may be administered to the patient. It may desirable to decrease or block expression of a particular IL4Ralpha isogene, which may be done by turning off by transforming a targeted organ, tissue or cell population with an expression vector that expresses high levels of untranslatable mRNA for the isogene. Specific therapeutics identified by these methods may be useful for allergic diseases. The present sequence is a probe for human IL4R-alpha.
                                                                                                                                                                                                                                                                       IGFBP2
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                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF46143 standard; DNA; 15 BP
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                                                                                                                                                                                                                                                                       oligonucleotide #982.
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nilarity 92.9%;
Conservative
                              condition
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                                 the retina;
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Pred. No. 2.4e+0
0; Mismatches
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2.4e+02;
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                                                                         Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see ARF45151 and ARF45151). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of
                                                                                                                                                                                                                                                         30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFB]-2 or IGFBB], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an an analysis and that inhibits or reduces growth factor mediated proliferation and/or inflammation -
                                                                                                                                                                                                                                                                                                                     AAF46144 standard;
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                                                                                                                                                                                                                        IGFBP2 oligonucleotide #983.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           skin disorders.
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                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hyperplasia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 12.4; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No.
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WO200078341-A1

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RESULT 446
AAF46266
ID AAF462
XX AAF462
XX AAF462
XX AAF462
XX AAF462
XX AAF462
XX AAF15e
KW Cytosh dt
KW IGF bi
KW IGF bi
KW Kerato
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Best Local :
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                                                                                                                                                                                                               Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see ARF45181 and ARF45181). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contracting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an antisement sequence is an antisement sequence.
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                      21-JUN-2000; 2000WO-AU00693
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                                                                                                                             Homo sapiens.
                                                                                                                                                               neovascular condition of the
                                                                                                                                                                              keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease;
hyperneovascular condition; hyperplasia; kidney disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             blood vessels or any other hyperplasia.
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Pred. No. 2.4e+(
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?.4e+02;
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                                                                     Query Match
Best Local S
Matches 13
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                                                                                                                                               Sequence 15
                                                                                                                                                                               blood vessels or any other hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          proliferation and/or inflammation
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                                                                                       Local Similarity
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CCGCAGCCAGCCGG 15
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                                                                                                                                               BP; 3 A; 7 C; 5 G; 0.U; 0 other;
                                                                       Conservative
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                                                                                       0.9%;
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Pred. No. 2.
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RESULT 447
AAA6467
ID AAF467
AC AAF467
AC AAF467
AC AAF467
AC AAF467
XX
OT 30-MAH
XX
Antise
KW Cytosis
KW Cytosis
KW IGF bi
KW OZOOO
KW IGF bi
KW 
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neovascular cond
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sapiens.

21-JUN-2000; 28-DEC-2000

2000WO-AU00693

99US-0140345.

(MURD-) MURDOCH CHILDRENS RES INST

Wraight

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RESULT 448
AAF49086/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide of the present invention (see AAF45151 and AAF45153-F45.61). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of
               WPI; 2001-041421/05
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Ameliorating the effects of a disorder, e.g. psoriasis, ladministering UV (ultra-violet) treatment (optional) and nucleic acid that inhibits or reduces growth factor media

mediated

cell

an antisense

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(MURD-) MURDOCH CHILDRENS

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Werther GA,

21-JUN-2000; 2000WO-AU00693

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neovascular

condition of the retina;

sapiens.

IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;

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                                                                                                                     nucleic acid that inhibits or reduces proliferation and/or inflammation -
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The present invention relates to a method

for ameliorating the effects

inhibiting

reducing growth

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contracting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of

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                                                                                                                                   Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an a nucleic acid that inhibits or reduces growth factor mediated
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RESULT 452
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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contracting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
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of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide of the present invention (see AAF45.51 and AAF45.53-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of tiskin, a hyperneovascular condition such as a neovascular condition of the state of the process of the p

skin, growth

factor-mediated malignancies,

of the

The Example

8; Page 85; 201pp;

English.

present invention relates to a method for ameliorating the effects

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RESULT 453
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keratosis, neoplasias,
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                                          CC The present invention relates to a method for ameliorating the effects CC of skin disorders. The method comprises contacting the skin with an CC antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 CC receptor, IGF binding protein [IGFBP]-2 or IGFBP3], which is capable of inhibiting or reducing growth factor mediated cell proliferation, CC inflammation and/or other disorders. The present sequence is an CC oligonucleotide which can be used to design the antisense CC oligonucleotide which can be used to design the antisense CC oligonucleotides of the present invention (see AAF45151 and CC AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, CC keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other CC sclerotic disease, kidney disease, hyperproliferation of the inside of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13;
                               otic disease, kidney disease, hyperversels or any other hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      بر
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                                                                                                                                                                                                                                                                                                        93; 201pp; English
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Pred. No. 2.
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.4e+02;
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Sequence 15

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Query Match Best Local

Similarity

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Conservative

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Mismatches

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Sequence

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G; 7 T; 0 other; Score 12.4; DB 1; Pred. No. 2.4e+02;

Matches

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                                                      The present invention relates to a method for ameliorating the effects CC of skin disorders. The method comprises contacting the skin with an CC antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 CC receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, CC inflammation and/or other disorders. The present sequence is an CC oligonucleotide which can be used to design the antisense CC oligonucleotides of the present invention (see AF45151 and AF45153-F45161). The method is useful for ameliorating the effects of CC psoriasis, inchhyosis, pityriasis, ruba, pilaris, serbornhoea, kaloids, CC keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the CC skin, a hyperneovascular condition such as a neovascular condition of the Skin, a hyperneovascular condition such as a neovascular condition of the CC sclerotic disease, kidney disease, hyperproliferation of the inside of CC blood vessels or any other hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ammliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense therapy, antiproliferative; antinflammatory, antipsoriatic; cytostatic; dermatological; cardiant; vorucide; ophthalmological; kelo skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pittyriasi IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris
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Pred. No. 2.
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s; pilaris;
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RESULT 456
AAC62187/c
The specification describes a method of synthesis of biologically active CC substances of determined structure directly in the cells of living crystalisms containing specific RNA or DNA sequence. The method is based CC on the hybridisation of two or more oligomers bound with biologically inactive substances to specific RNA or DNA in vivo in the cells of CI inactive substances to specific RNA or DNA in vivo in the cells of CI inactive substances to specific RNA or DNA in vivo in the cells of CI inactive precursors bound to the oligomers can interact with each other CC inactive precursors bound to the oligomers can interact with each other CC in the continuous and reactions which bind the biologically inactive form of the substances. This changing of properties CI is due to chemical reactions which bind the biologically active form of the whole compound. The methods are useful for producing biologically continuous from inactive precursors. These compounds may be considered to the compounds from inactive precursors. These compounds may be continuous from inactive precursors. These compounds may be continuous from the cellular metabolism. DNA replication, RNA cranscription, RNA translation, RNA elongation RNA processing, protein continuous processing, cellular metabolism. DNA replication, CCI in channel transmission, cellular protein and RNA transportation, CCI processes of cellular oxidation, toxins, proteins or RNAs. Oligomers are CC Ancastalel to human NbHOT, which is specific to ovarian cancer cells. The method of the invention is used to produce the tumour suppressor p53, and the oligomers are constant of the control of the foundation of the specific to ovarian cancer cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Biologically active compound; cellular metabolism; DNA replication; RNA transcription; RNA translation; RNA clongation; RNA processing; protein synthesis; protein processing; cellular differentiation; cell division; ion channel transmission; cellular protein; toxin; RNA transportation; cellular oxidation; tumour suppressor p53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             producing proteins or polynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel methods for the synthesis of biologically active compounds from inactive precursors in the cells of living organisms, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-006911/01.
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                                      from
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                                    bound peptides and
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                                      oligomers.
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Query Match Best Local Similarity

0.9**%**; 92.9**%**;

Score Pred. w T; 0

12.4; DB 1; No. 2.4e+02; other;

Length 15;

Query Match Best Local

y Match 0.9%; Local Similarity 92.9%;

Conservative

<u>,,</u>

Score 12.4; DB 1; Pred. No. 2.4e+02; Mismatches

Length Indels

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Sequence

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RESULT 457
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                                                                                            This invention relates to a novel method for preventing phagocytosis of immune complexes comprises introducing an inhibitor of a kinase endogenous to phagocytic cells associated with an Fc receptor at the membrane of the cells under conditions so that the phagocytic potential of the cells is inhibited. The method of the invention may have immunosuppressive, dermatological, antiinflammatory, antiarthritic, antirheumatic and antiasthmatic activities and may be used as a kinase inhibitor. The method and compositions of the invention may be used for modulating the clearance of antibody coated cells, viruses and soluble antigens by inhibiting phagocytosis and modulating the interaction of immune complexes with cellular to tissue Fc receptors. The method is used for treating autoimmune diseases, immune mediated diseases e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-SEP-1998;
30-SEP-1993;
30-SEP-1994;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                        Preventing phagocytosis of immune complexes used for treating autoimmune diseases comprises introducing inhibitor of kinase endogenous to phagocytic cells associated with Fc receptor at of cells -
                           rheumatoid arthritis, and for preventing immune complexes deposition tissues e.g. the kidneys and in the joints. The present sequence represents a human Syk kinase gene target III antisense oligonucleotiused in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 phagocytosis; immune complex; kinase inhibitor; immune mediated disease; asthma; systemic '''''''' rheumatoid arthur''
                                                                                                                                                                                                                                                                                            Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Syk mRNA target III antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US2002068703-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rheumatoid arthritis; antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABX15428
                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYPE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E 1649 TGCTGGCAGGGGTC 1662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13;
                                                                                   and immune complex diseases e.g. lupus erythematosus and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ss; Syk; kinase; immunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         VINU
                                                                                                                                                                                                                                                                                            6; Fig
15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGCTGGCAGGGATC
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B₽;
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                                                                                                                                                                                                                                                                                           9; 26pp;
0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                              Park J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93US-0129381
94US-0316425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96US-0657884
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-0158980
3 C; 3 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                           English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        dermatological;
                                         antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antiasthmatic;
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                                                                                                                                                                                                                                                                                                                                                                  e.g.
                                                                                                                                                                                                                   potential
                                                                                                                                                                                                                                                                                                                                         membrane
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RESULT 459
AAT60192/c
ID AAT601
XX
AC AAT601
XX
DT 03-FEB
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AAD49046,
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                                                                                                                                                                                   Query Match
Best Local S
Matches 13
                                                                                                                                                                                                                                                                 The present invention relates to novel peptide modulators of androgen receptor. The peptides of the invention are useful for treating androgen-associated disorders such as prostate cancer, particularly hormonally refractive prostate cancer, colon cancer, lung cancer, benign prostatic hypertrophy, acne, hirsutism, male pattern baldness, Stein-Leventhal syndrome, androgen insensitivity syndrome, infertility, endometrial cancer and X-linked spinal bulbar muscular atrophy. The present DNA sequence is MVDP gene androgen responsive element (ARE).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Androgen receptor; androgen-associated disorder; prostate cancer; acne; benign prostatic hypertrophy; hirsultism, androgen insensitivity syndrome; male pattern baldness; Stein-Leventhal syndrome; infertility; cytostatic; X-linked spinal bulbar muscular atrophy; antiseborrheic; dermatological; depilatory; androgen responsive element; ARE; MVDP gene; ds.
                             AAT60192;
 03-FEB-1998
                                                         AAT60192 standard;
                                                                                                                                                                                                                                           Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 26; 68pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                         New peptide modulators of androgen receptor, useful for treating androgen-associated disorder, e.g. prostate cancer, particularly hormonally refractive prostate cancer, colon cancer, lung cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-MAR-2001; 2001US-275240P.
28-JAN-2002; 2002US-352399P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-MAR-2002; 2002WO-US07487.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200272612-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD49046;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD49046 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PRAE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1783
                                                                                                                                                       2633 GAAGTTCTTGTTCT 2646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2003-067363/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene androgen responsive element (ARE).
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                                                                                                                                                                                     l Similarity
13; Conservat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GACAAAGACAAGCC 1796
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                                                                                                                              GAAGTTCCTGTTCT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mueller J,
                                                                                                                                                                                                                                          BP; 2
                                                                                                                                                                                     Conservative
 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                                                                           A; 3 C; 3 G; 7 T; 0
                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oza VB,
                                                         16
                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                  Score 12.4; DB 1;
Pred. No. 2.4e+02;
0; Mismatches 1;
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                                                                                                                                                                                                                                             other;
                                                                                                                                                                                                              Length 15;
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                                                                                                                                                                                     Gaps
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RESULT 460
AAA86559/c
ID AAA865
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents a ribozyme recognition site for the proliferating cell nuclear antigen (PCNA) gene which is cleaved by a harpin ribozyme at position 867 and by a hammerhead ribozyme at position 869. Novel ribozymes are being investigated for their ability to inhibit the activity of a growth factor (e.g. PCNA) responsible for abnormal smooth muscle cell (SMC) proliferation in vascular tissue leading to restenosis. The ribozymes can also directly block the production of oncogenes and cell regulatory factors involved with SMC growth following vascular injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New hairpin and hammer:head ribozyme(s) - which inhibit abnormal smooth muscle cell proliferation in vascular tissue, partic. for preventing or treating restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ribozyme; hairpin; hammerhead; proliferating cell nuclear antigen; growth factor; oncogene; vascular tissue; SMC; PCNA; recognition site; restenosis; smooth muscle cell proliferation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
             04-DEC-1998;
                                      06-DEC-1999;
                                                                                                                                                          Ribozyme; hairpin; hammerhead; gene therapy; vasotropic;
                                                                                                                                                                                                                                                                 AAA86559 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 16 BP; 4 A; 4 C; 1 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 15; 50pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-SEP-1996;
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                                                                08-JUN-2000
                                                                                         WO200032765-A2
                                                                                                                                             restenosis;
                                                                                                                                                                                  PCNA hairpin ribozyme recognition site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                        2169 TTTGGTAACAGAAA 2182
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                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                              (first entry)
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             98US-0110954.
                                       99WO-US28772.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96WO-US14838.
                                                                                                                                                                                                                                                                                                                                                                                                0.9%;
92.9%;
                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                  Score 12.4; DB 1;
Pred. No. 2.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                               Length 16;
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RESULT 461
AAA86780/c
ID AAA867
XX AAA867
XX AAA867
XX PCNA h
XX Ribozy
KW R
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent lother than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given it AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells.
                                                                      The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDKI, PCNA and Cyclin B1.
                       Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for
                                                                                                                                                                               Example
                                                                                                                                                                                                                       New hairpin and hammerhead ribozyme RNA encoding a cyclin or cell-cycle PCNA and Cyclin B1 -
                                                                                                                                                                                                                                                                                                                                WPI; 2000-412314/35
                                                                                                                                                                                                                                                                                                                                                                                  Tritz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                    (-DWMI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-JUN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200032765-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mammalia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ribozyme; hairpin; hammerhead; gene therapy; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCNA hammerhead ribozyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 16 BP; 4 A; 4 C; 1 G; 7 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           efficient
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tritz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (-DWMI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoding a cyclin or cell-cycle and Cyclin B1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ribozyme is resistant to endonuclease activity and cient in restenosis treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2169 TTTGGTAACAGAAA 2182
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                                                                                                                                                                                                                                                                                                                                                                                                                                    IMMUSOL INC.
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                                                                                                                                                                          1; Page 24; 109pp; English
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                                                                                                                                                                                                                                                                                                                                                                            Welch PJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0110954
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%;
                                                                                                                                                                                                                                                                                                                                                                                  Barber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Barber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   recognitoin site #5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
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                                                                                                                                                                                                                                                                                                                                                                               дŖ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred.
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                                                                                                                                                                                                                                                    for inhibiting restenosis, dependent kinase other than
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for inhibiting restenosis, cleaves dependent kinase other than CDK1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.6e+02
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                   The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatiis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent
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13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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Pred. No. 2.
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Best Local :
ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antistickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; wulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP;
                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 394; 408pp; English.
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Pred. No. 2.
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RESULT 464
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                            method. The method comprises obtaining a restriction fragment length polymorphism (RFLP) pattern of the 24 rpoB gene fragments; isolating, amplifying and digesting the DNA fragment from the microorganism to be identified and comparing the RFLP patterns from the known rpoB gene fragments with the unidentified fragment. The rpoB gene fragments are useful to identify a wide range of Mycobacterium species, e.g. for diagnosis or to obtain epidemiological and pathogenesis information for selection of appropriate therapies, including M. tuberculosis, M. leprae and non-tuberculous mycobacteria (NTM) encountered in subjects infected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH57577 to AAH62099 represent sequences used in the
                                                                                                                                                                                                                                                                                                           ligonucleotide probes (AAS05227-AAS05242) that can be used to detect specific mycobacterial species. The probes are described in an invention relating to the use of rpoB gene fragments (AAS05201-AAS05224) from various Mycobacterial species. These rpoB gene fragments can be used in the diagnosis and identification of Mycobacterium species using a novel PCR restriction fragment length polymorphism analysis (PRA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mycobacterium kansasii.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Non-tuberculous mycobacteria; rpoB gene fragment; NTM; HIV; PRA; RFLP; PCR-restriction fragment length polymorphism analysis; probe; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                restriction fragment length polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New DNA fragments from the rpoB gene of mycobacteria, useful for diagnosis and identification of many mycobacterial species by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 16 BP; 4 A; 4 C; 1 G; 7 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present sequence for Mycobacterium kansasii oligonucleotide pe KANSASII can be used to detect M. kansasii. It is 1 of 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ë
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Pred. No. 2
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human immunodeficiency virus

(HIV).

Analysis of the

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RESULT 465
ABN84079/c
cengineered to contain 2 template annealing sequences, i.e. a
complate, e.g. mRNA encoding at least a portion of an antibody, and
cengiate, e.g. mRNA encoding at least a portion of an antibody, and
complate, e.g. mRNA encoding at least a portion of an antibody, and
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complate encoding at least a portion site located
complate encoding sequence. The collar sequence is
complated to encoding and and
complated to encoding an encoding sequence (see ABN84080), their junction
complated to encoding an encoding sequence and esparated by a restriction digest to
complate a cleaved plasmid. The portion of the cleaved plasmid that
complated to the antibody constant region serves as a primer for
complated transcriptase to generate first strand cDNA. Transcription
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                                                                                                                                                                                                                                                                   in situ production of genes. It was found nucleic acids encoding at least a portion of an antibody can be directly incorporated into a plasmid by reverse transcription of mRNA. The plasmid is
                                                                                                                                                                                                                                                                                                                   The present sequence is an antibody framework region 1 (FR1) collar oligonuclectide that is used in the engineering of plasmids of the invention, especially plasmid pRL5-CAT (see ABN84078), for the
                                                                                                                                                                                                                                                                                                                                                                                                                                           Plasmid for in situ production of genes, comprises two template annealing sequences, such as downstream primer and upstream collar sequence and a restriction site located between the annealing sequences
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required), and can dif
experiment, including
                                                                                                                                                                                                                                                                                                                                                                                            Claim 35;
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                                                                                                                                                                                                                                                                                                                                                                                          Page 18; 65pp; English
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                                                                                                                     Novel sitosterolemia susceptibility gene polypeptide and polynucleotide, useful for screening a compound that increases the level of expression or activity of SSG polypeptide for treating
                                                                                                               level of expression or a sterol-related disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; sitosterolaemia susceptibility gene; SSG; atherosclerosis; sterol-related disorder; hyperlipidaemia; hypercholesterolaemia; thera gall stone; coronary heart disease; cardiovascular disease; arthritis;
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                                                                                                                                                                                                                                                                                                                                                 misc_feature
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                                                                                                                                                                                                                                                                                    25-OCT-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                   xanthoma; haemolytic
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                                                                                                                                                                   2002-017598/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sitosterolaemia
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                                                                                                                                                                                      Schultz J,
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2000US-204234P.
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                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; 2 C; 6 G; 5 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                    anaemia; transgenic animal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       susceptibility gene (SSG)
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"Exon"
                                                                                                                                                                                                                                                                                                                                                           "Intron"
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The invention relates to an isolated Sitosterolaemia Susceptibility Gene (SSG) polypeptide. SSG is a member of adenosine triphosphate (ATP) binding cassette (ABC) family cholesterol transporter. SSG is useful for identifying a compound useful in the treatment or prevention of a sterol-related disorder, including sitosterolaemia, hyperlipidaemia, hypercholesterolaemia, gall stones, HDL deficiency, atherosclerosis or hypercholesterolaemia, gall stones, HDL deficiency, atherosclerosis or

Fig 14B; 105pp; English.

hypercholesterolaemia, gall nutritional deficiencies. S

associated diseases or conditions including

SSG is also useful for treating

coronary heart

cholesterol-

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RESULT 467
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                                                                                                                                                          The invention relates to a novel isolated molecule comprising bases 2-8 or 13-16 of 2 16 base pair sequences, or comprising a 1731 base pair sequence, all given in the specification or at least 95 % identity with the 1731 bp sequence. The nucleic acid molecule is useful in regulating apoptosis in cells and in drug screening. The method is useful in facilitating the induction of apoptosis in cells, in identifying an agen that can facilitate the induction of apoptosis in cells, and in inhibiting the growth of a cancer. This polynucleotide sequence represents a ribozyme binding substrate sequence relating to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             grug
                                                                                                                                                                                                                                                                                                                                                                                                                                                               and
                                                                                              Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 42; 153pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated nucleic acid molecule useful for regulating apoptosis induction in cells, for inhibiting the growth of cancer in subjects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-129308/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tritz R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ribozyme substrate binding
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                                                                                            A; 3 C; 3 G; 3 T; 0 other;
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Score 12.4; DB 1;
Pred. No. 2.6e+02;
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Pred. No. 2.6e+02;
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                                                                                                                                                patient previously diagnosed with a breast tumour and subsequently useful in determining the degree of aggressive treatment indicated. The prognostic method is especially useful in making decisions concerning treatment modalities, including therapeutic intervention, diagnostic criteria such as disease staging and disease monitoring and surveillance for metastasis or recurrence of neoplastic disease. The present sequence is a PCR primer used for LOH analysis. This primer is used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                          tumour re-occurrence in a patient previously diagnosed with a breast tumour. The method involves analysing the target cell sample for loss of heterozygesity (LOH) to chromosomal locus 3p24.3. The method is useful for determining the likelihood of tumour re-occurrence in a
                                                                                                                                                                                                                                                                                                                                                                                               Determining the likelihood of tumor re-occurrence in a patient previously diagnosed with a breast tumor to determine treatment modalities, comprises analyzing the cell sample for loss of heterozygosity to chromosomal locus 3p24.3 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified.
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                                                                                                                       Sequence 16 BP; 2 A; 5 C; 4 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                        The invention relates
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dairkee SH,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-MAR-2002; 2002WO-US09068.
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                                                            <u>,</u>
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Pred. No. 2.6e+02;
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                   19-1631-717A-5
18-678-039A-14
17-947-683-14
17-947-683-13
18-471-570-17
19-014-241-13
19-277-078-13
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COMPTTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA RE-LEASE #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/471,570
FILING DATE: 06-JUN-1995
CLASSIFICATION DATN:
APPLICATION DATN:
APPLICATION NUMBER: US/08/149,664
FILING DATE: 16-AUG-1991
APPLICATION NUMBER: US 07/743369
FILING DATE: 16-AUG-1991
APPLICATION NUMBER: US 08/14369
FILING DATE: 16-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: LINEK, Ernest V
REGISTRATION NUMBER: 29822
REFERENCE/DOCKET NUMBER: 40897
TELECOMUNICATION INFORMATION:
TELEPHONE: (617)523-3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: DAVID G. CONLIN;
ADDRESSEE: CUSHMAN
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US-09-064-156A-301
US-09-064-156A-302
US-09-071-845-242
US-09-071-845-547
US-09-071-845-547
US-09-071-845-547
US-09-071-845-664
US-09-071-845-664
US-09-071-845-730
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418, App
445, Appl
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638, App
639, App
640, App
1246, Ap
2267, Ap
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Best Local Similarity
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APPLICATION NUMBER: US/07/642,755
FILING DATE: 18-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Oblon, No. 5288855man F.
REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 769-226-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)521-4500
                                                                                                                                                                                                                                              TELEPHONE: (703)521-45
TELEFAX: (703)486-2347
TELEX: 248855 OPAT UR
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Extracellular Form of the Human TITLE OF INVENTION: Fibroblast Growth Factor Receptor
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APPLICANT: Mazue,
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                                                                                                                                                           STRANDEDNESS: uni
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CITY: Arlington
STATE: Virginia
ZIP: 22202
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                                                                  Conservative
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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Pred. No. 2
                                                                Mismatches
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                                                               Gaps
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US-08-471-570-15/c Sequence 15, Patent No. 57 GENERAL INFO

INFORMATION:

CITY: Boston STREET:

130 Water Street

Massachusetts

02109

TELEFAX: (617)523-6440
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO:

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RESULT 4
US-07-631-717A-5/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; MOLECULE TYPE: DNA (genomic) US-07-997-133-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 769-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)521-4500
TELEFAX: (703)486-2347
TELEFAX: (703)486-2347
TELEX: 24885 OPAT UR
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 1.8%;
Best Local Similarity 93.1%;
                                                                                                                                                                                       GENERAL
                                                                                           APPLICANT: Yayon, Avner
APPLICANT: Ornitz, David M.
APPLICANT: Klagsbrun, Michael
APPLICANT: Leder, Philip
APPLICANT: Leder, Philip
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/07/642,755
FILING DATE: 18-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Oblon, Norman F.
REGISTRATION NUMBER: 24,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/997,133
FILING DATE: 28-DEC-1992
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Bergonzoni, APPLICANT: Mazue, Guy
                                        TITLE OF INVENTION: SYSTEM FOR ASSAYING BINDING TITLE OF INVENTION: TO A HEPARIN-BINDING GROWTH TITLE OF INVENTION: FACTOR RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sarmientos, Paolo
TITLE OF INVENTION: Extracellular Form of the Human
TITLE OF INVENTION: Fibroblast Growth Factor Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: VII
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Arlington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                          2488 TGTTGGCATGCAGTGCCCTCCCAGAGACC 2516
                                                                                                                                                                                     INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isacchi, Antonella
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Pred. No. 2.
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US-08-166-717D-5/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APHILING DATE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Paul T. Clark
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00383/018001
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 10542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence 5, Application US/08166717D Patent No. 5789182
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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APPLICANT: Yayon, Avner
APPLICANT: Ornitz, David M.
APPLICANT: Klagsbrun, Michael
APPLICANT: Leder, Philip
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (617) 542-89
TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION: I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
COMPUTER: IBM PS/2 Model SC
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WordPerfect (Version 7.0)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/166,717D
FILING DATE: 12/14/93
CLASSIFICATION: 435
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PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 19901220
                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PS/2 model 50Z or 55S OPERATING SYSTEM: IBM P.C. DOS (Ver SOPTWARE: WordPerfect (Version 5.0)
                                                                                                                                                               COUNTRY: UZIP: 02110
                                                                                                                                                                                                     STATE:
                                                                                                                                                                                                                                    ADDRESSEE:
STREET: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 1.6%;
Local Similarity 95.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1873 GAGATGGAGATGAAGATGATT 1896
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                                                                                                                                                                                                                       Boston
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                                                                                                                                                                                                 Massachusetts
                                                                                                                                                                                                                                    E: Clark & Elbing LLP
176 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   225 Franklin
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IBM P.C. DOS (Version 3.30)
                                                                                                                                                                                                                                                                                                           SYSTEM FOR ASSAYING BINDING TO A HEPARIN-BINDING GROWTH FACTOR RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US/07/631,717A
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Pred. No. 8.4;
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US-08-678-039A-4/c
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                                   US-08-678-039A-4
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 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                     TELEFAX: 202-783-6031
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                 MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Saxe, Stephen A.
REGISTRATION NUMBER: 38,609
REFERENCE/DOCKET NUMBER: 23
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Kristina Bieker-Brady
REGISTRATION NUMBER: 39,109
REFERENCE/DOCKET NUMBER: 00383/017002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 723-4123
TELEFAX: (617) 723-8962
                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Keating, APPLICANT: Morris,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 12/20/90
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/678,039A FILING DATE: 10-JUL-1996 CLASSIFICATION: 435
                                                                                     TOPOLOGY: linear
                                                                                                   STRANDEDNESS: single
                                                                                                                                                                                                         TELEPHONE: 202-624-1589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 1.6%;
Local Similarity 95.8%;
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                                                                                                                                     LENGTH:
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                                                                                                                     H: 25 base pairs nucleic acid
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555 Thirteenth Street, N.W., Suite 701 East
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tower
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                                                   /desc = "Primer se
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Diagnosis of Williams Syndrome and Williams Syndrome Cognitive Profile by Analysis of the Presence or Absence of a LIM-Kinase Gene
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Score 21.4;
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Pred. No. 8.
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Length 25;

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US-07-947-683-14
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                                                                                              Sequence 13, Application US/08400323
Patent No. 5703047
GENERAL INFORMATION:
APPLICANT: wilson, Steven E.
TITLE OF INVENTION: Methods and OTHER OF INVENTION: Healing with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 14, Applicat Patent No. 5589451 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                      Query Match 1.5%;
Best Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
Matches 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 33,928
REFERENCE/DOCKET NUMBER: UTSD:311/KIT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 512-320-7200
TELEPHONE: 512-474-7577
TELEX: NOT APPLICABLE
INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/07/947,60
FILING DATE: SEPTEMBER 21, 1992
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: KITCHELL, BARBARA S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPB: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY
                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                              ADDRESSEE: Arnold, White & Durkee STREET: P. O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U
ZIP: 77210
                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: HOUSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
                                                                                                                                                                                                                                                                                               1322 TATCCTTTCACTCTGCATGGT 1342
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                                                                                                                                                                                                                                                                      21 TATCCTTTCACTCTGCATGGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14, Application US/07947683
o. 5589451
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                                                                                                                                                                                                                                                                                                                                          21;
Houston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WILSON, STEVEN E.

VENTION: METHODS AND TREATMENTS FOR

VENTION: CORNEAL HEALING WITH HEPATOCYTE
                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                              Methods and Treatments for Corneal Healing with Growth Factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AND KERATINOCYTE GROWTH FACTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WHITE & DURKEE
                                                                                                                                                                                                                                                                                                                                      1.5%; Score 21; DB
100.0%; Pred. No. 7.8
1ve 0; Mismatches
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0; Mismatches
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COUNTRY: ZIP: 772

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US-08-471-570-16
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Patent No. 5
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: IGAKASDA, APPLICANT: IGAKASDA, APPLICANT: SENOO, MABAHATU
APPLICANT: WATANABE, Tatsuya
APPLICANT: WATANABE, TATSUYA
APPLICANT TAVENTION: PROTEIN, DNA AND USE THEREOF
                           FILING DATE:
FILING DATE:
APPLICATION NUMBER: US 07/743369
FILING DATE: 16-AUG-1991
ATTORNEY/ACENT INFORMATION:
NAME: LINEK, Ernest V
REGISTRATION NUMBER: 29822
REGISTRATION NUMBER: 29822
                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/149,664
                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
TELEPAX: (512) 474-7577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
FELECOMMUNICATION, INFORMATION:
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FILING DATE: 09-MAR-1995
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                       APPLICATION NUMBER: US/01
FILING DATE: 06-JUN-1995
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ZIP: 02109
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TOPOLOGY: lir
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                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 1.5%; Score 21; DB Local Similarity 100.0%; Pred. No. 7. es 21; Conservative 0; Mismatches
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5750371
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                   40897
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RESULT 10
US-08-471-570-17/c
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Best Local Similarity
Matches 21; Conserv
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Query Match 1.4%;
Best Local Similarity 84.6%;
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                     TELEX: 200291 STRE UR INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 07/
FILING DATE: 16-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: LINEK, Ernest V
REGISTRATION NUMBER: 29822
REFERENCE/DOCKET NUMBER: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: other nucleic acid, synthetic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 06-JUN-
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/471,570
                                                                                    MOLECULE TYPE: other nucleic acid, synthetic DNI
                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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APPLICANT: WATANABE, Tatsuya
TATE OF TNVENTION: PROTEIN, DNA AND USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: SENOO,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                        TELEPHONE: (617)523-6440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U
ZIP: 02109
                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE:
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TELEX: 200291 STRE UR
                                                                                                     TOPOLOGY:
                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Boston
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STRANDEDNESS: single
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                                                                                                                                     nucleic acid
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                                                                                                                                                       26 base pairs
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                                                                                                       linear
                                                                                                                                                                                                                                       (617)523-3400
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                                                                                                                     single
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                                                                                                                                                                                                                                                                          40897
                   Score 19.6;
Pred. No. 22;
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                                   Length 26;
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0; Mismatches

Indels

0

Gaps

1388 CCCCAGACTACCTGGAGATAGCCATT 1413

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APPLICANT: ANDERSSON, Leif
APPLICANT: MOLLER, Maria J
APPLICANT: MOLLER, Maria J
APPLICANT: WALLES, Richard
APPLICANT: SIGGENS, Renneth W.
APPLICANT: SIGGENS, Kenneth W.
APPLICANT: PLASTOW, Graham S.
TITLE OF INVENTION: Methods for Determining the Coat Colour Genotype of a
TITLE OF INVENTION: Pig
FILE REFERENCE: 064727.0103
CURRENT APPLICATION NUMBER: US/09/014,241
CURRENT FILING DATE: 1998-01-27
EARLIER APPLICATION NUMBER: PCT/GB96/01794
EBARLIER APPLICATION NUMBER: GB 9515385.4
EARLIER APPLICATION NUMBER: GB 9515385.4
EARLIER APPLICATION NUMBER: GB 9525364.7
EARLIER APPLICATION NUMBER: GB 9525364.7
EARLIER FILING DATE: 1995-12-12
NUMBER OF SEQ ID NOS: 16
SOPTWARE: Patentin Ver. 2.0 - beta
SEQ ID NO 13
LENGTH: 22
Type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Description of Artificial Sequence: Derivative ; OTHER INFORMATION: pig sequence US-09-014-241-13
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US-09-014-241-13
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Best Local Similarity 90.9
                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 39, Application US/09277078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -09-v1.
Sequence 13, Approximate No. 6183955
                                                                                                                                                                                                                                     APPLICANT: Sakurada, Kazuhiro
APPLICANT: Palmer, Theo
APPLICANT: Palmer, Theo
APPLICANT: Cage, Fred H.
TITLE OF INVENTION: REGULATION OF TYROSINE HYDROXYLASE
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: 07251/031001
CURRENT APPLICATION NUMBER: US/09/277,078
CURRENT FILING DATE: 1999-03-26
                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 6312949
GENERAL INFORMATION:
                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                   PEATURE:
NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: n =
                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                 OTHER INFORMATION: Oligonucleotide for PCR
                                                                                                                                                                  ENGTH: 20
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Pred. No. 21;
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RESULT 14 US-07-741-940-61/c

Sequence 61, Application US/07741940 Patent No. 5352775
GENERAL INFORMATION:

APPLICANT:

ALBERTSEN, HANS

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                                                                                                           ; HYPOTHETICAL:
; ANTI-SENSE: 1
US-08-951-923-33
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US-08-951-923-33/c
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                                                       Matches
                                                                  Query Match
Best Local Similarity
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Best Local Similarity
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Ve:

CURRENT APPLICATION DATE: US/08/951,923

FILING DATE: October 16, 1997
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TELEX: 380816COOLEYPA
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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LOCATION: (0)...(0)
                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Neeley, Richard L.
REGISTRATION NUMBER: 30,092
REFERENCE/DOCKET NUMBER: BITT-001/02US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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20 CAGACTTTGGACTAGCCAGA 1
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VENTION: PHENOTYPIC ASSAYS OF CYCLIN/CYCLIN-DEPENDENT KINASE
                                                     Conservative
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Pred. No. 19
                                                                  Score 18.4;
Pred. No. 2
                                                       Mismatches
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US-08-289-548A-61/c
; Sequence 61, Application US/08289548A
; Patent No. 5648212
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NAME: Kagan, Sarah A.

REGISTRATION NUMBER: 32,141

REFERENCE/DOCKET NUMBER: 1107.035574

TELECOMMUNICATION INFORMATION:

TELEPHONE: 202-508-9100

TELEPAX: 202-508-9299

INFORMATION FOR SEQ ID NO: 61:

SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                 APPLICANT:
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ORIGINAL SOURCE:
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APPLICANT: WARKHAM, ALEXANDER F.
APPLICANT: NAKAMURA, YUSUKE
APPLICANT: THLIVERIS, ANDREW
TITLE OF INVENTION: INHERITED AND SOMATIC MUTATIONS OF APC
TITLE OF INVENTION: GENE IN COLORECTAL CANCER IN HUMANS
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                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                       INVENTION:
                                                               GRODEN, JOANNA
HEDGE, PHILIP J.
JOSLYN, GEOFF
KINZLER, KENNETH
MARKHAM, ALEXANDER F.
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1001 G Street, NW
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NAKAMURA, YUSUKE
THLIVERIS, ANDREW
UENTION: INHERITED AND SOMATIC MUTATIONS OF APC
VENTION: GENE IN COLORECTAL CANCER IN HUMANS
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HEDGE, PHILIP J.
                                                                                                                                             CARLSON, MARY
GRODEN, JOANNA
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                                                                                                                                                                                                    ALBERTSEN, HANS
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87.0%; Pred. No. 32;
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 61, Application US/08452654
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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APPLICANT: ALBERT
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: FORM TYPE
MEDIUM TYPE: FORM TYPE
MEDIUM TYPE
M
                            ZIP: 20001-4598
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner, Birc
STREET: 1001 G Street,
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ADDRESSEE: Banner & Allegretti,
STREET: 1001 G Street, NW
                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: INHERITED AND SOMATIC MUTATIONS OF APC
TITLE OF INVENTION: GENE IN COLORECTAL CANCER IN HUMANS
NUMBER OF SEQUENCES: 94
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TELECOMMUNICATION INFORMATION:
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Local Similarity 87.0%;
es 20; Conservation
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STRANDEDNESS: sing
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HEDGE, PHILIP J.
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THLIVERIS, ANDREW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KINZLER, KENNETH MARKHAM, ALEXANDER F.
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GRODEN, JOANNA
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PatentIn Release #1.0,
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reet, NW
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Pred. No. 3;
                                                                                                                                                                                                                                                                                                                                              McKie & Beckett
   Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         32;
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                                                    COUNTRY: USA

ZIP: 20001-4598

ZIP: 20001-4598

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OBERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNUMBER: US/08/452,655B
FILING DATE: 25-MAY-1995
CLASSIFICATION 530
PRIOR APPLICATION DATA:
NEW TOWN NUMBER: US/08/452,655B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 08-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 1107
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
TELEFAX: 202-508-9299
INFORMATION FOR SEQ ID NO: 61:
SEQUENCE CHARACTERISTICS:
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APPLICANT:
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APPLICANT:
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ORIGINAL SOURCE:
ORGANISM: Hom
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                APPLICATION NUMBER: US 00 FILING DATE: 12-AUG-1994 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: INHERITED AND SOMATIC MUTATIONS OF APC
TITLE OF INVENTION: GENE IN COLORECTAL CANCER IN HUMANS
NUMBER OF SEQUENCES: 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
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CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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Local Similarity 87.0%;
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GRODEN, JOANNA
HEDGE, PHILIP J.
JOSLYN, GEOPF
KINZLER, KENNETH
MARKHAM, ALEXANDER F.
NAKAMURA, YUSUKE
THLIVERIS, ANDREW
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1001 G Street, NW
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linear
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                                                  US 08/289,548
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US 07/741,940
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Pred. No. 32;
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RESULT 18
US-08-450-582-61/c
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Best Local Similarity 87.0%;
Matches 20; Conservative
                                                                                                                                                               COUNTRY: USA
ZIP: 20001-4598
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPACTION
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/450,582
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No.
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APPLICANT: ALBERT
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                                                             CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US 08/452,655
FILING DATE: 25-MAX-1995
APPLICATION NUMBER: US 08/289,548
FILING DATE: 12-AUG-1994
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APPLICANT:
APPLICANT:
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ATTORNEY/AGENT INFORMATION:
                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/741,940
FILING DATE: 08-AUG-1991
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
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                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                STATE: D.C
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STREET: 10
                                                                                                                                                                                                                                                                                                                                               CITY: Washington
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM:
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MARKHAM, ALEXANDER F.
NAKAMURA, YUSUKE
THLIVERIS, ANDREW
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1001 G Street, NW
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CARLSON, MARY
GRODEN, JOANNA
HEDGE, PHILIP J.
JOSLYN, GEOFF
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                                                                                                                                                                                                                                                                                                                                                                                                                                   INHERITED AND SOMATIC MUTATIONS OF APC
GENE IN COLORECTAL CANCER IN HUMANS
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Pred. No. 3;
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US-08-450-582-61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 1.3%;
Best Local Similarity 87.0%;
Matches 20; Conservative
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SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
INFORMATION FOR SEQ ID NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
ORIGINAL SOURCE:
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                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/449,731
FILING DATE: 25-May-1995
CLASSIFICATION CUNKNOWN>
PRIOR APPLICATION WMBER: 08/289,548
FILING DATE: 12-AUG-1994
ATTORNEY AGENT INFORMATION:
                                                                            REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION: TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS_DOS
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ANAND, RAKESH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: INHERITED AND SOMATIC MUTATIONS OF APC
GENE IN COLORECTAL CANCER IN HUMANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
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                                                                                                                                      NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Washington
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                                                             TELEFAX: 202-508-9299
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HEDGE, PHILIP J.
JOSLYN, GEOFF
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THLIVERIS, ANDREW
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MARKHAM, ALEXANDER F.
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Pred. No. 32;
                                                                                                                      1107.46943
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RESULT 21
                                                                                                                                                                 ; ANTI-SENSE: US-08-951-923-26
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US-08-951-923-26/c
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Patent No. 6048693
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                                                                                                         Matches
                                                                                                                                    Query Match
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Ve.

CURRENT APPLICATION NUMBER: US/08/951,923

FILING DATE: October 16, 1997

FILING DATE: October 16, 1997
                                                                                                                                                                                                                                                                                                       TELEFAX: 650 857-0663
TELEX: 380816COOLEYPA
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                               MOLECULE TYPE: HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                      TYPE: STRANDEDNESS: SING
                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                NAME: Neeley, Richard L.
REGISTRATION NUMBER: 30,092
REFERENCE/DOCKET NUMBER: BITT-001/02US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 87.0
                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
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STREET: 5 Palo Alto Square, 3000 El Camino Real
                                                                                                                                                                                                                                                                          ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: cDNA ORIGINAL SOURCE:
                                                                       2203 GACTTTGGACTCGCCAGAGAT 2223
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                                                                                                        l Similarity
19; Conserv
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                                                                                                                                                                                                                                                           nucleic acid
                                            GACTTTGGACTAGCCAGAGCT 1
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650 857-0663
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                                                                                                      Score 17.8; D
Pred. No. 28;
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Pred. No. 32;
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US-08-910-629A-62/c

Sequence 62, Application US/08910629A Patent No. 5877309

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APPLICANT: McKay, Robert A.

APPLICANT: Dean, Nicholas M.

APPLICANT: Monia, Brett

APPLICANT: Nonia, Brett

APPLICANT: Nonia, Brett

APPLICANT: Nonia, Brett

TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS

FILE REFERENCE: ISPH-0350

CURRENT APPLICATION NUMBER: U$/09/287,796A

CURRENT FILING DATE: 1999-04-07

EARLIER APPLICATION NUMBER: 09/130,616

EARLIER APPLICATION NUMBER: 08/910,629

EARLIER APPLICATION NUMBER: 08/910,629

EARLIER APPLICATION NUMBER: 1997-08-03
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                                                                                                                                                                                                                                                            Sequence 62, Application US/09287796A
Patent No. 6133246
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                                                                                                                                                                                                                                            ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Jane Massey Licata
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (609) 779-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (609) 779-841
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: Line
ANTI-SENSE: Yes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: WINDOWS SOFTWARE: WORDPERFECT 6.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE,
MEDIUM TYPE: STORAGE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: Nucleic Acid
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Brett Monia
                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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COMPOSITONS AND METHODS FOR THE MODULATION OF JNK
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APPLICANT: Sakurada, Kazuhiro

APPLICANT: Palmer, Theo

APPLICANT: Palmer, Theo

APPLICANT: Palmer, Theo

ITITLE OF INVENTION: REGULATION OF TYROSINE HYDROXYLASE

ITITLE OF INVENTION: EXPRESSION

FILLS REFERENCE: 07251/031001

CURRENT APPLICATION NUMBER: US/09/277,078

CURRENT APPLICATION NUMBER: US/09/277,078

CURRENT FILLING DATE: 1999-03-26

NUMBER OF SEQ ID NOS: 60

SOFTWARE: FRATSEQ for Windows Version 4.0

SEQ ID NO 40

LENGTH: 20
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US-09-277-078-40/c
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US-09-130-616-62/c
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SEQ ID NO 62
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                               Sequence 40, Application US/09277078 Patent No. 6312949
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local Similarity
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LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: McKay, Robert A.
APPLICANT: Dean, Nicholas M.
APPLICANT: Monia, Brett
APPLICANT: Nero, Pam
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLECTIDE COMPOSITI
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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CURRENT APPLICATION NUMBER: US/09/130,616C
CURRENT FILING DATE: 1998-08-07
EARLIER APPLICATION NUMBER: 08/910,629
EARLIER FILING DATE: 1997-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 62, Appli
Patent No. 6221850
OTHER INFORMATION: Oligonucleotide for PCR
                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
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                     FEATURE
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nes 17; Conserva
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100.0%; Pred. No.
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100.0%; Pred. No.
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35;
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                                                                                         US-09-433-699-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Donna T. Ward
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF RECQL2 EXPRESSION
FILE REFERENCE: RTS-0207
CURRENT FILING DATE: 2001-03-01
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 38
SEQ ID NO 38
SEQ ID NO 38
CORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 38, Applicate Patent No. 6399378 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                      Query Match
Best Local Similarity
                                                                                                                                                                        SEQ ID NO 42
                                                                                                                                                                                                                                                                                                    Sequence 42, Application US/09433699B
Patent No. 6165786
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Best Local Similarity
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                                                                                                                                                                                 APPLICANT: C. Frank Bennett
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF NUCLEOLIN EXPRESSION
FILE REFERENCE: RTS-0109
CURRENT APPLICATION NUMBER: US/09/433,699B
CURRENT FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 89
                                                                                                                             LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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NAME/KEY: misc_feature
LOCATION: (0)...(0)
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OTHER INFORMATION: h = A,
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NAME/KEY: misc_feature
                                                                                                   FEATURE:
OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
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hes 17; Conserv
1878, GGAGATGATGAAGATGAT 1895
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                               Conservative
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                                           Score 16.4;
Pred. No. 49
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Pred. No. 38;
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Pred. No. 3
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US-08-863-639A-65; Sequence 65, Application US/08863639A; Patent No. 5981185
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US-08-863-639A-44/c
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                                                                                                                                                                              Patent No. 5981185
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08,
FILING DATE: May 28, 1997
CLASSIFICATION: 435
                                                                                     APPLICANT: Matson, Robert S.
APPLICANT: Coassin, Peter J.
APPLICANT: Rampal, Jang B.
APPLICANT: Caskey, C. T.
TITLE OF INVENTION: OLIGONUCLEOTIDE REPEAT ARRAYS
NUMBER OF SEQUENCES: 95
                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Matson, Robert APPLICANT: Coassin, Peter
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                                                                        CORRESPONDENCE ADDRESS:
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REGISTRATION NUMBER:
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COUNTRY:
                                          STREET:
                                                          ADDRESSEE:
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es 18; Conserv
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Pred. No. 53;
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CURRENT APPLICATION NUMBER: US/09/100,398; CURRENT FILING DATE: 1998-06-19; NUMBER OF SEQ ID NOS: 17; SOFTWARE: PATENTIN VEr. 2.0; SEQ ID NO 2; LENGTH: 20; TYPE: DNA; ORGANISM: primer
US-09-100-398-2
                                                                                                                       RESULT 30
US-09-226-012-39/c
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US-09-100-398-2/c
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GENERAL INFORMATION:
APPLICANT: Keating, Mark T.
APPLICANT: Splawski, Igor
APPLICANT: Splawski, Igor
TITLE OF INVENTION: MUTATIONS IN AND GENOMIC STRUCTURE OF HERG
TITLE OF INVENTION: SYNDROME GENE
                                                                                      Sequence 39, Application US/09226012
Patent No. 6207383
                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Conrad, Danie APPLICANT: Kelly, Ann E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (626) 795-632: INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: JOSEPH E. Mueth
REGISTRATION NUMBER: 20,532
REFERENCE/DOCKET NUMBER: 1185
TELECOMMUNICATION INFORMATION:
TELEPHONE: (626) 796-4000
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TOPOLOGY: linear
MOLECULE TYPE: Oth
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Local Similarity 89.5%;
es 17; Conservation
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SOPTWARE: Corel WordPerfect 8 version
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N: 435
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795-6321
NO: 65:
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; Pred. No. 53;
0; Mismatches
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Pred. No. 5
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US-08-584-040-7597; Sequence 7597, A
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US-09-733-294A-61
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Best Local S
Matches 17
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LENGTH: 20
TYPE: DNA
ORGANISM: Homo sapiens
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LENGTH: 20
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Best Local Similarity 89.5%;
Matches 17; Conservative
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                                                                                                                                                                                                                    Patent No. 6346398
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: SUSAN M. Freier
APPLICANT: Edward V. Wancewicz
TITLE OF INVENTION: ANTISENSE MODULATION OF TERT EXPRESSION
FILE REFERENCE: ISPH-0527
CURRENT APPLICATION NUMBER: US/09/733,294A
CURRENT FILING DATE: 2000-12-07
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CURRENT APPLICATION NUMBER: US/09/226,012
CURRENT FILING DATE: 1999-01-06
EARLIER APPLICATION NUMBER: 09/122,847
EARLIER FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 09/572,423
PRIOR FILING DATE: 2000-05-16
NUMBER OF SEQ ID NOS: 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Brett P.
                                                              TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                    APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, Ja
APPLICANT: Stinchcomb, D
                              CORRESPONDENCE ADDRESS:
                                                   NUMBER OF SEQUENCES:
                                                                                                                                                        APPLICANT:
                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                           2020 GGGATGGAGTACTCCTATG 2038
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633 West Fifth Street
                                                                                                                                                                    McSwiggen, James
Stinchcomb, Dan
                                                                                                                                                                                                                                                        Application US/08584040
                                                                                                                                                   Escobedo,
                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                 METHOD AND REAGENT FOR THE
TREATMENT OF DISEASES OR
CONDITIONS RELATED TO LEVELS
OF VASCULAR ENDOTHELIAL
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                                                                 GROWTH FACTOR
                                                                                                                                                        Jaime
                                                                                                                                                                    Dan T.
                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 56;
0; Mismatches
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; LENGTH: 17
; TYPE: RNA
; ORGANISM: Mus sp.
US-09-371-772B-3391
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                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3391, Application US/09371772B Patent No. 6566127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                    Query Match
                                                                                                                                                                                                                                                                                  APPLICANT: PAVCO, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MeBHB00, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam APPLICANT: McSwiggen, Jim APPLICANT: Stinchcomb, Dan APPLICANT: Escobedo, Jaime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
PILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U$/08/584,040
FILING DATE: January 11, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
/ Match 1.1%;
Local Similarity 76.5%;
nes 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Com
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
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Similarity 76.5%;
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  3; Mismatches
                    Score 15.4;
Pred. No. 47;
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Pred. No. 4
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                                        DB 1;
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                                    Length 17;
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  Gaps
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RESULT 35
US-09-555-889A-5/c
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.0 SEQ ID NO 4818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4818, Application US/09371772B Patent No. 6566127
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Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Escobedo, Jaime

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

FILE REFERENCE: MBHB00,876-J (237/198)

CURRENT FILING DATE: 1990-08-10

PRIOR APPLICATION NUMBER: US 60/005,974

PRIOR APPLICATION NUMBER: US 60/005,974

PRIOR FILING DATE: 1995-10-26

PRIOR PILING DATE: 1995-10-26

PRIOR FILING DATE: 1996-01-08

NUMBER OF SEO ID NOS: 14225

COUNTAINE DESCRIPT VERSION 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam APPLICANT: McSwiggen, Jim APPLICANT: Stinchcomb, Dan APPLICANT: Escobedo, Jaime
                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOPTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION UMBER: US/09/555,889A

FILING DATE: 09-Apr-2001

INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Nucleotide sequence encoding the tomato light hypersensitive phenotype, coded proteins and uses thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Bowler, Chris
Mustilli, Anna Chiara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2105 CCAGAGGCATGGAGTAC
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LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Robert J. Jondle
STREET: 555 13th Street NW,
CITY: Washington
STATE: District of Columbia
                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
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Pred. No. 47;
                                                                                                                                                                                                                                                                                                                                                                                              Suite 701-E
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US-08-913-050A-3

### Sequence 3, Application US/08913050A

### Patent No. 5827726
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US-08-368-704C-89/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; SEQUENCE DESCRIPTION: SEQ ID NO: 5: US-09-555-889A-5
                                                                                                                                                                                                                  US-08-368-704C-89
                                                                                                                                                       Query Match
Tocal Similarity
                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                         TELEFAX: (202) 371-254
TELEX: 248636 SSK
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/080,850
PILING DATE: 24-UN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/258,287
FILING DATE: 10-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Yuan, Junying APPLICANT: Miura, Masayuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                  NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.3920002
                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/368,704C FILING DATE: 4-JAN-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TLE OF INVENTION:
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                                                                                                                                                                                                                                                                            ENGTH:
                                                                                                               1878-GGAGATGATGAAGATGA 1894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1816 GCCGTGAAGATGTTGAA 1832
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                                                                                                                                                                                                                                                           nucleic acid
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                                                                                                                                                                                                                                                                           20 base pairs
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1100 New York Avenue,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kessler, Goldstein & Fox ork Avenue, Suite 600
                                                                                                                                                                  Score 15.4;
Pred. No. 6
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                                                                                                                                                    Pred. No. 66;
); Mismatches
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Pred. No. 59
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                                                                                                                                                                               Length 20;
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; OTHER INFORMATION: Antisense Oligonucleotide US-09-658-688A-24
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                                                                                                                                                                                                                                                                                                                                      RESULT 38
US-09-658-688A-24
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                                                                   NUMBER OF SEQ ID NOS:
SEQ ID NO 24
LENGTH: 20
TYPE: DNA
                                                                                                                                                                                                                            GENERAL INFORMATION:

APPLICANT: Donna T. Ward
APPLICANT: William Gaarde
APPLICANT: Brett P. Monia
                                                                                                                                                                                                                                                                                                  Sequence 24, Application US/09658688A Patent No. 6498035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 1.1%;
Best Local Similarity 70.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: JP 57104/1995
PILING DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP PCT/JP96/00660
PILING DATE: 15-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: YUN, Allen C.
REGISTRATION NUMBER: 37,971
REGISTRATION NUMBER: 37,971
                                                                                                                                       TITLE OF INVENTION: ANTISENSE MODULATION OF MEKK3 EXPRESSION FILE REFERENCE: RTS-0143 CURRENT APPLICATION NUMBER: US/09/658,688A CURRENT FILING DATE: 2000-09-08
                                                                                                                                                                                           APPLICANT: Jacqueline Wyatt TITLE OF INVENTION: ANTIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (202) 737-3528
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: DNA ENCODING PROTEIN KINASE NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEB: BROWDY ****
STREET.
                                 ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/913,050A
FILING DATE: 05-SEP-1997
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 20004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Washington STATE: D.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                             1813 GTGGCCGTGAAGATGTTGAA 1832
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                                                                                                                                                                                                                                                                                                                                                                                                          nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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Pred. No. 7
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Query Match

1.1%; Score 15.2;

DB 1;

Length 20;

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RESULT 40
US-09-944-036-19
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US-09-198-452A-5279
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APPLICANT: Griffals, R.
APPLICANT: Griffals, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preve
TITLE OF INVENTION: and treatment of infection
FILE REFERENCE: 9710-003-99
CURRENT REPERENCE: 9710-003-99
CURRENT FILING DATE: 1998-11-24
RUMBER OF SEQ ID NOS: 6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19, App. -
Sequence 19, App. -
No. 6582920
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LENGTH: 20
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  Best Local Similarity
                           Query Match
                                                                NAME/KEY: modified base
LOCATION: (1)..(2)
OTHER INFORMATION: 2'-O-methyladenosine
NAME/KEY: modified base
LOCATION: (3)..(4)
OTHER INFORMATION: gm
-09-944-036-19
                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: BABOLA, Odile
APPLICANT: TRAN, Nathalie
APPLICANT: VENUT, Guy
APPLICANT: VENUT, Guy
TITLE OF INVENTION: AMPLIFICATION OF HIV-1 SEQUENCES FOR DETECTION OF
TITLE OF INVENTION: SEQUENCES ASSOCIATED WITH DRUG-RESISTANCE MUTATIONS
FILE REFERENCE: GP114-02 UT
CURRENT APPLICATION NUMBER: US/09/944,036
CURRENT FILING DATE: 2001-08-31
PRIOR APPLICATION NUMBER: US 60/229,790
PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 70
CORPURABE OF SEQ ID NOS: 70
CORPURABE OF SEQ ID NOS: 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: YANG, APPLICANT: BREN' APPLICANT: BABO!
                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                            LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Chlamydia pneumoniae
                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Oligonucleotide primer for Protease target OTHER INFORMATION: sequence
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Local Similarity 85.0%;
es 17; Conservation
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  1.1%;
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Pred. No. 71;
Score 15.2; DB 1;
Pred. No. 71;
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RESULT 42
US-08-585-684B-1806
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                                                                                                                                         Matches
                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                             TELEFAX: (213) 955-044
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FASTSEQ VERSION 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,68
FILING DATE: January 16, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/000,951
FILING DATE: JULY 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: WARDURG, RICHARD
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIE: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
COMPUTER: TOWN D C DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 633 mer.
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                1.1%;
Local Similarity 73.3%;
les 11; Conservative
                                                                                                                                                                                                                                            TYPE: nucleic STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A. ZIP: 90071
                                                                                                                                                                                                                                 TOPOLOGY:
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                                                                                                                                                                                                                                                                                  LENGTH:
                                                                                                      1373 AGGAGATTACAGCTT 1387
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                                                                     AGGAGAUUACAGCUU 15
                                                                                                                                                                                                                                                                                15 base pairs
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633 West Fifth Street
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                                                                                                                                                                                                                                                                                                                                                      (213) 955-0440
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                                                                                                                                                           Score 15; DB
Pred. No. 42;
                                                                                                                                           Mismatches
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Sequence 1806, Application US/08585684B Patent No. 5877021

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GENERAL INFORMATION: APPLICANT: Stinch

APPLICANT:

Jarvis, Thale McSwiggen, Ja

.METHOD AND REAGENT FOR THE

Stinchcomb, Daniel

FITLE OF INVENTION:

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US-08-585-684B-1807
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                                                                                                                                                                                                                                                                                                                           Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local Similarity 73.3%;
                                                                                                                                                                                                                                                                                                                                         sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/585,684B
FILING DATE: January 16, 1996
PRIOR APPLICATION DATA:
APPLICATION UMBER: 60/000,951
FILING DATE: July 7, 1995
ATTORNEY AGENT INFORMATION:
NAME: WARDLUTG, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 67-3510
NFORMATION FOR SEQ ID NO:
                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                 TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE INDUCTION OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES NUMBER OF SEQUENCES: 2751
                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ELECURATE (213)
                                                                                                                                                                                                                                                                                            PPLICANT:
COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.(
                                                                                                         CITY: Los Angeles
STATE: California
                                                                                                                                         STREET:
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                                                                                         COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 1373 AGGAGATTAÇAGÇTT 1387
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California
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                                                                                                                                                    E: Lyon & Lyon
633 West Fifth Street
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633 West Fifth Street
                                                                                                                                         Suite 4700
                                                                                                                                                                                                                                                              McSwiggen,
                                                                                         U.S.A.
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                                                                                                                                                                                                                                                                                         Stinchcomb, Daniel T.
                                                                                                                                                                                                                                                                           Jarvis, Thale
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FastSEQ Version 1.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      955-0440
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                                                                                                                                                                                                                                                              James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1806:
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Pred. No.
 DOS 5.0
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                                              1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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APPLICANT: Stinch
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/(
FILING DATE;
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/;
NFORMATION FOR SEQ ID NO: 1805:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Watburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: January 16, 1996 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
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                                                   TELEPHONE: (213)
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COMPUTER: 150.

OPERATING SYSTEM: IBM F...

OPERATING SYSTEM: IBM F...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1374 GGAGATTACAGCTTC 1388
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>. 6194150
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                                  67-3510
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               633 West Fifth Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             U.S.A.
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                                                                955-0440
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                                                                                                                                                                                                                                                                                                                                 08/585,684
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DOS 5.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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US-09-038-073-1806
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US-09-038-073-1806
                                                                       Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 6194150
GENERAL INFORMATI
                                                                                                                                                                                      TELEFAX: (213) 955-044
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: IBM P.C. DO:
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Jarvis, Thale APPLICANT: McSwiggen, James
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                    TELEPHONE: (213) 955-0440
                                                                                                                                        TYPE: nucleac
STRANDEDNESS:
                                                                                                                                                                                                                                                                                    NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 633 West Fifth Street
STREET: Suite 4700
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                                                                      Local
                                                                                                                                                          LENGTH: 15 base pairs
TYPE: nucleic acid
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                             1373 AGGAGATTACAGCTT 1387
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ll Similarity 73.3%;
11; Conservative
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nucleic acid
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AGGAGAUUACAGCUU 15
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                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   IBM Compatible
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                                                                                                                                                                                                                                               (213) 489-1600
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                                                                      1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       METHOD AND REAGENT FOR THE INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES 2751
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                                                                                                                                                                                                                                                                                                                                                  08/585,684
                                                                                                                                                                                                      1806:
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                                                                       Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                     DOS 5.0
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                                                                                   DB 1; Length 15;
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US-08-951-923-51/c
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US-09-038-073-1807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1807, App
Patent No. 6194150
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                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                  Sequence 51, Application US/08951923
Patent No. 6048693
                                                                                                                                                                     GENERAL INFORMATION: APPLICANT: Bitter
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APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                NUMBER OF SEQUENCES: 5
                                                                                                                                 APPLICANT: Bitter, Grant TITLE OF INVENTION: PHEN TITLE OF INVENTION: FUNC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
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APPLICATION NUMBER: US
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Local Similarity 73.3%;
les 11; Conservativo
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OPERATING SYSTEM:
                                 CITY: Palo Alto
STATE: CA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 633 West F. STREET: Suite 4700
ZIP: 94306-2155
                 COUNTRY:
                                                                  STREET:
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                                                                                  ADDRESSEE:
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: California
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                                                                  5 Palo Alto Square, 3000 El Camino Real
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633 West Fifth Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FastSEQ Version 1.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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                                                                                  Cooley Godward LLP
                                                                                                                   PHENOTYPIC ASSAYS OF CYCLIN/CYCLIN-DEPENDENT KINASE FUNCTION
57
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                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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RESULT 49
US-09-632-580A-65/c
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US-08-951-923-51
                                            Patent No. 6255111
GENERAL INFORMATION:
                                                               Sequence 65, Application US/09632580A Patent No. 6255111
                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 87
SEQ ID NO 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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Best Local Similarity
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TITLE OF INVENTION: ANTISENSE MODULATION OF G-ALPHA-16 EXPRESSION
FILE REFERENCE: RTS-0032
CURRENT APPLICATION NUMBER: US/09/205,143
CURRENT FILLING DATE: 1998-12-03
APPLICANT: C.
APPLICANT: Lex M.
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 650 857-0663
TELEX: 380816COOLEYPA
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                              LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                  FEATURE:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
ANDLICATION NUMBER. US (06/05/102)
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TELEPHONE: 650 843-5000
TELEFAX: 650 857-0663
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: OCCLASSIFICATION:
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16; Conserv
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               Cowsert
                               Frank Bennett
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ANTISENSE MODULATION OF HER-4 EXPRESSION
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Pred. No. 67;
0; Mismatches
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RESULT 50
US-08-149-105-10
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Best Local Similarity 88.9%;
Matches 16; Conservative
Matches
                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 65
LENGTH: 18
                                                                                                                                                                                                                        APPLICATION NUMBER: 08/029,673
FILING DATE: March 11, 1993
APPLICATION NUMBER: 07/853,396
FILING DATE: March 18, 1992
ATTORNEY/AGENT IMFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00786/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: RTS-0054
CURRENT APPLICATION NUMBER: US/09/632,580A
CURRENT FILING DATE: 2000-07-31
NUMBER OF SEQ ID NOS: 93
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                                                                                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Antisense Oligonucleotide
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APPLICATION NUMBER: US/08/149,105
                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 530 PRIOR APPLICATION DATA:
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MEDIUM TYPE: 3.5" Di
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ADDRESSEE: Fish & Richardson
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                                                                        STRANDEDNESS:
TOPOLOGY: lin
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Local Similarity
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                                                                                                                                                                               (617) 542-8906
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Conservative
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SYSTEM: MS-DOS (Version 5.0)
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            Score 14.6;
Pred. No. 6
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Pred. No. 6
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Mismatches
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                          DB 1;
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                         Length 17;
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Indels
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US-08-317-847-10
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                                                                                                                                      US-08-529-878B-20/c
                                                                                                                                                            RESULT 52
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                                                                          Sequence 20, Application US/08529878B Patent No. 5932556 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/029,673
FILING DATE: March 11, 1993
APPLICATION NUMBER: 07/853,396
FILING DATE: March 18, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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APPLICANT: Gustafson
APPLICANT: He, Wei W
TITLE OF INVENTION:
TITLE OF INVENTION:
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OPERATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/317,847
APPLICANT: Tam, Robert C.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TITLE OF INVENTION: REGULATION OF CD28 EXPRESSION NUMBER OF SEQUENCES: 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
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MEDIUM TYPE: 3.5" Diskette,
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ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
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                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FAMILY
                                                                                                                                                                                                                                                                                            1.0%; Score 14.6; I
82.4%; Pred. No. 64;
tive 2; Mismatches
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US-08-541-950B-17
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Patent No. 5821046
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/541,950B
FILLING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/960,370
FILLING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M.
REGISTRATION NUMBER: 34,380
REGISTRATION NUMBER: 3777/57347
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9110
TELEPHONE: (617) 345-9110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 714-525-3433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Fish, Robert D.
REGISTRATION NUMBER: 33,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WOO'DEPAFECT 6.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall CTITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                      CIT:. Mr. STATE: Mr. 02111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 13-SEI CLASSIFICATION: 424
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California
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                                                                                                                                                                                                                                                                                                   Diskette, 3.50 inch, 1.44 Mb storage
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                                                            3777/57347 (MRC-011AX)
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Pred. No. 63
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US-08-541-950B-20
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                                                                      Query Match
Best Local Similarity
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                                                          Matches
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                                                                                                                                                                                                                                                                                TELEFAX: (617) 345-91:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/541
FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/960,37
FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathle
REGISTRATION NUMBER: 34,380
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
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ADDRESSEE: Banner & Witcoff, Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C
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                                                                                                                             NAME/KEY: misc_feature
LOCATION: 8
OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Diskette, 3.50 inch, COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                           STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Boston
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OTHER INFORMATION: N is 2'-deoxythymidine
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                          AGCCAGACTTCAGCAGC 1506
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AGCCAGANUUGAGCAGC 17
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                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                             Kathleen M.
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Pred. No. 69
                                                                        Score 14.4;
Pred. No. 69;
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                                                        Mismatches
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                                                                                    DB 1;
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                                                                                   Length 17;
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RESULT 56
US-09-083-756A-20
; Sequence 20, Application US/09083756A
; Patent No. 6114109
; GENERAL INFORMATION:
; APPLICANT: KAIT J, Gait MJ, Heaphy :
; TITLE OF INVENTION: VIRAL (HIV) GROWNER OF SEQUENCES: 26
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Best Local Similarity
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (617) 345-91
INFORMATION FOR SEQ ID NO:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatibl
OPERATING SYSTEM: PC-DOS/N
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
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NAME: Williams, Ph.D., K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                     CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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                                              CILL.
STATE: MA
STATE: 02111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
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ZIP: 02111
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                                                                                        CITY: Boston
                                                                                                        STREET:
                                                                                                        ADDRESSEE: Banner & Witcoff, Ltd.
STREET: One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: 8 OTHER INFORMATION: N is 2'-deoxythymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature 
LOCATION: 8
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REFERENCE/DOCKET NUMBER: 3777/57347 (MRC-Ollax)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/083,756A
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                                                                                                                                                                                                                                                                                                                                                  1490 AGCCAGACTTCAGCAGC 1506
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One Financial Center, 45th Floor
                                                                                                                                                                      Karn J, Gait MJ, Heaphy S, Dingwall C
                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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Pred. No. 6
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; OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine US-09-083-756A-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-584-040-5715
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY_AGENT INFORMATION:
ATTORNEY_AGENT INFORMATION:
NAME: Williams, Ph.D., Kathle
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 3777
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (617) 345-9111
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 17 bases
                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF
TITLE OF
                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                PRIOR APPLICATION DATA:
                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WOLECULE TYPE: synthetic RNA
                                                                                                                                                                                                                                                                                                                                             UMBER OF SEQUENCES:
                                               FILING DATE: Jackson CLASSIFICATION:
                                                             APPLICATION NUMBER: US/08/584,040 FILING DATE: January 11, 1996
                                                                                                            OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                STATE:
                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/083,756A
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                                                                                                                                                                                                                              COUNTRY:
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                                                                                                                                                                                                                                                                                                            ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1490 AGCCAGACTTCAGCAGC 1506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION:
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13; Conserv
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                                                                                                                                                                                                                                             Los Angeles
: California
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                                                                                                                                                                                                                                                                                                                                                                                                                          INVENTION:
                                                                                                                                                                                                                                                                                            E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                               Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/08584040
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                                                                                                                                            IBM Compatible
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60/005,974
r 26, 1995
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                                                                                                                               DOS 5.0
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US-09-371-772B-2598
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Best Local Similarity
Matches 12; Conserv
                              Sequence 23, Application US/08541950B Patent No. 5821046 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.0 SEQ ID NO 2598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 08/584,040 PRIOR FILING DATE: 1996-01-08 NUMBER OF SEQ ID NOS: 14225
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor FILE REFERNCE: MEHBOO, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PPLICANT:
                                                                                                                                                                                                                                                                                                                                                      ENGTH:
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TITLE OF INVENTION:
                  APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 218/064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX:
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                                                                                                                                                                                           1822 AAGATGTTGAAAGATG 1837
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EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pavco, Pam
McSwiggen, Jim
Stinchcomb, Dan
Escobedo, Jaime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17 base pairs
Karn J, Gait MJ, Heaphy S, Dingwall C
WENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (213) 955-0440
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                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                              Score 14.4; D
Pred. No. 69;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ა
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Pred. No. 69;
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                                                                                                                                                                                                                                                               Length 17;
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NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:

ADDRESSEE: STREET: O

One Financial Center, 45th Floor

Banner & Witcoff, Ltd

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RESULT 61
US-09-083-756A-23
; Sequence 23, Apj
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US-09-205-922-30
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                                                                                                                                                                                                                                                             SEQ ID NO 30
LENGTH: 18
TYPE: DNA
                                                                                                                             Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Lex M. COMBERT
TITLE OF INVENTION: ANTISENSE MODULATION OF G-APLHA-11 EXPRESSION
FILE REFERENCE: RTS-0030
CURRENT FILING NUMBER: US/09/205,922
CURRENT FILING DATE: 1998-12-04
NUMBER OF SEQ ID NOS: 87
                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 30, Appli
Patent No. 5951455
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                            OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 18 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/960,370
FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: WordPerfect
CURRENT APPLICATION DATA
APPLICATION NUMBER: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
mes 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: 8
OTHER INFORMATION: N is 4-thio-2'-deoxythymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Williams, Ph.D., REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: PC-DOS/N
SOFTWARE: WordPerfect 6.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
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                                                                                               1814 TGGCCGTGAAGATGTT 1829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1490 AGCCAGACTTCAGCAGC 1506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleic acid
                                                                            TGGCGGTGAAGATGTT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGCCAGANUUGAGCAGC 17
 Application US/09083756A
                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/09205922
                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             synthetic RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diskette, 3.50 inch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
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                                                                                                                                                   Score 14.4;
Pred. No. 78;
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Pred. No. 78
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                                                                                                                                        Mismatches
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                                                                                                                                                                 DB 1;
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                                                                                                                                                                 Length 18;
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                                                                                                                                     Indels
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RESULT 62
US-09-658-645A-5
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US-09-658-645A-5
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                                                                                                   SEQ ID NO 5
                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity Matches 13; Conserv
                                                                                                                                                                                                                                                                                                Sequence 5, Application US/09658645A Patent No. 6423518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                   TITLE OF INVENTION: Design and Production of Mutant 2,5-Diketo-D-gluconic TITLE OF INVENTION: Acid Reductase Enzymes with Altered Cofactor Dependen FILE REFERENCE: RU-0078
CURRENT APPLICATION NUMBER: US/09/658,645A
CURRENT FILING DATE: 2000-09-11
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                             APPLICANT: Anderson, Stephen APPLICANT: Banta, Scott
                                                                                                                    SOFTWARE: PatentIn Ver.
                                          ORGANISM: Artificial Sequence
                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (617) 345-91:
            OTHER INFORMATION: Description of Artificial Sequence: Synthetic
                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/083,756A FILING DATE: PRIOR APPLICATION DATA: APPLICATION NUMBER: 08/541,950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: synthetic RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOPTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C
TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: N is 4-thio-2'-deoxythymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Williams, Ph.I
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ZIP: 02111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Banner & Witcoff, Ltd.
STREET: One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                           1490 AGCCAGACTTCAGCAGC 1506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid
DEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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Pred. No. 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-371-772B-4817
                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/338,907
CURRENT FILING DATE: 1999-06-23
EARLIER APPLICATION NUMBER: 08/996,306
EARLIER FILING DATE: 1997-12-22
EARLIER APPLICATION NUMBER: 60/099,658
EARLIER FILING DATE: 1998-09-09
EARLIER APPLICATION NUMBER: 09/218,207
EARLIER FILING DATE: 1998-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.0 SEQ ID NO 4817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 84, Application US/09338907 Patent No. 6265546
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Best Local Similarity 85.7%;
Matches 12; Conservative
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CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
CURRENT FILING DATE: 1999-08-10
CRICK FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-28
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                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Ilya, Chumakov
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: PROSTATE CANCER GENE
FILE REFERENCE: GENSET.18CP1CP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENERAL INFORMATION:
                                                                                                                                                                       NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cohen, Daniel
APPLICANT: Blumenfeld, Marta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 14225
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                                                          TYPE: DNA
ORGANISM: Mus Musculus
AME/KEY: misc feature
                                                                                                          ENGTH: 17
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Escobedo, Jaime
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McSwiggen, Jim
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llarity 93.8%;
Conservative
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Pred. No. 81;
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0; Mismatches
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US-08-584-040-4205
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US-09-218-207-84/c
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LENGTH: 17
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Best Local Similarity
Matches 15; Conser
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                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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SOFTWARE: Patent.pm
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EARLIER FILING DATE: 1997-12-22
EARLIER APPLICATION NUMBER: 60/099,658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: GENSET.018CP1
CURRENT APPLICATION NUMBER: US/09/218,207
CURRENT FILING DATE: 1998-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Cohen, Daniel
APPLICANT: Blumenfeld, Marta
APPLICANT: Ilya, Chumakov
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: Prostate cancer gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EARLIER FILING DATE: 1998-09-09
                                                                                                                            TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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LOCATION: 1..17
OTHER INFORMATION: sequencing oligonucleotide moPGrace5R444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Mus Musculus
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                                                                                                                                                                                                          APPLICANT: Stinchcomb
APPLICANT: Escobedo,
                                                                                                                                                                                                                                          APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                             NUMBER OF SEQUENCES:
                                             STREET: 633 West Fifth Street STREET: Suite 4700
COUNTRY:
                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 1.0%;
Local Similarity 88.2%;
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                              Los Angeles
                                                                                                                                                                                                                                                                                              6346398
             California
                                           Suite 4700
                                                                                                                                                                                                             Stinchcomb, Dan T. Escobedo, Jaime
                                                                                                                                                                                                                                                                                                            Application US/08584040
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                                                                             Lyon & Lyon
                                                                                                                                             METHOD AND REAGENT FOR THE
TREATMENT OF DISEASES OR
CONDITIONS RELATED TO LEVEL
OF VASCULAR ENDOTHELIAL
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                                                                                                                                GROWTH FACTOR
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Pred. No. 87;
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Pred. No. 87;
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APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TOTAL OF INVENTION: OF VASCULAR ENDOTHELIAL
TOTAL OF INVENTION: CONDITIONS RELATED TO LEVELS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-584-040-4205
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Best Local S
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INFORMATION FOR SEO ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 17 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                  SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: January 1
                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/0
FILING DATE: October 26,
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS
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                                                                                                            COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: Januar CLASSIFICATION: 514
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                                  CLASSIFICATION:
                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2112 CATGGAGTACTTGGCTT 2128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 CAUGGAGUUCUUGGCAU 17
                                                                                                                                                                                                           90071-2066
                                                                                                                                                                                                                                                                                              E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                Suite 4700
                                                                                                                                                                                                                               U.S.A.
                                                                                                   IBM Compatible SYSTEM: IBM P.C. Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/08584040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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January 11, 1996
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                                                UMBER: US/08/584,040
January 11, 1996
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60/005,974
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Pred. No. 8
                                                                                                                     DOS 5.0
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RESULT 68
US-08-584-040-4242
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SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.0%;
Best Local Similarity 58.8%;
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No. 6346398
GENERAL INFORMATION:
                                                                       CLASSIFICATION: 514

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974

PILING DATB: October 26, 1995

ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327

REFERENCE/DOCKET NUMBER: 218/064

TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,0
FILING DATE: January 11, 1996
                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Pavco, rumnames
APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.

**Grobedo, Jaime
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: METHOD AND RÉAGENT FOR THE TITLE OF INVENTION: TREATMENT OF DISEASES OR TITLE OF INVENTION: CONDITIONS RELATED TO LEVEI TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL TITLE OF INVENTION: GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                   TELEFAX: (21)
TELEFAX: 67-3510
TELEFAX: 67-3510
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                                   TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2113 ATGGAGTACTTGGCTTC 2129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 AUGGAGUUCUUGGCAUC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                           90071-2066
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633 West Fifth Street
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Pred. No. 8
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US-08-584-040-4357
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Best Local
                                                         Matches
                                                                      Query Match
Best Local Similarity
                                                                                                                                                                        TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                                                                                                                   NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA: APPLICATION NUMBER: 60/005,974
                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                          TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                 STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                           TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: Januar
                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Con
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.0%;
Local Similarity 52.9%;
es 9; Conservation
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                                                                                                                                                                                                                                                                                                                                            FILING DATE: October 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2323 AGTGATGTCTGGTCCTT 2339
                          2404 GAACTTTTTAAGCTGCT 2420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INVENTION: METHOD AND REAGENT FOR THE INVENTION: TREATMENT OF DISEASES OR INVENTION: CONDITIONS RELATED TO LEVELS INVENTION: OF VASCULAR ENDOTHELIAL INVENTION: GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGUGACGUCUGGUCUUU 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   McSwiggen, James
Stinchcomb, Dan T.
Escobedo, Jaime
                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pavco, Pamela
                                                                                                                                                                                                                                                                                                                                                                                                                                              Word Perfect 5.1
                                                                                                                                 linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                storage
                                                                                                                                             single
                                                                      1.0%;
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Y 11, 1996
                                                      Score 13.8; D
Pred. No. 87;
6; Mismatches
                                                                                                                                                                                                                                                                                    218/064
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Pred. No. 8
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                                                                                  Length 17;
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RESULT 71 US-08-584-040-5779

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Sequence 5779, Application US/08584040
Patent No. 6346398
GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Stinchomb, Dan T.
APPLICANT: Stinchomb, Dan T.
APPLICANT: Bscobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE

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                                                                                                                                US-08-584-040-5714
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US-08-584-040-5714
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Patent No. 6346398
                                                                Matches
                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATB: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                            NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCS/DOCKET NUMBER: 216
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/005,97 FILING DATE: October 26, 1995 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.
ZIP: 90071-2066
                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Los Angeles
                                                                                                                                                                                                                                                              TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                1813 GTGGCCGTGAAGATGTT 1829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION:
CANT: Pavco, Pamela
                                                                10;
1 GUAGCCGUCAAGAUGUU 17
                                                                                                                                                                                                                                                  67-3510
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633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             McSwiggen, James
Stinchcomb, Dan
                                                                                                                                                                                                                                                                  (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Escobedo,
                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/08584040
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                                                                                                                                                 linear
                                                                              1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            METHOD AND REAGENT FOR THE TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GROWTH FACTOR
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                                                                                                                                                                                                                                                                                                                218/064
                                                                                Score 13.8;
Pred. No. 8
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                                                                  Mismatches
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                                                                                               Length 17;
                                                                  Indels
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RESULT 72
US-08-584-040-5780
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                                                                                                                                                                                                                                                                                           Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                 APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: October 26, ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                   TITLE OF INVENTION:
                                                                                                 TUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
TOPOLOGY: li
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/005,974
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                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                             2112 CATGGAGTACTTGGCTT 2128
                                                                                                                                                                                                                                                                                   1 5780, Apr.
                                                                                                                                                                                                                                                                       INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
10; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90071-2066
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Los Angeles
California
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                                                E: Lyon & Lyon
633 West Fifth Street
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633 West Fifth Street
                                 Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Suite 4700
                                                                                                                                                                                                                                                         Pavco,
                                                                                                                                                                                                                                                                                                           Application US/08584040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                         Pamela
                                                                                                                                 TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL
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                                                                                                   GROWTH FACTOR
8502
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 13.8; | Pred. No. 87; | S; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: OCCODER 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/06
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                  APPLICANT: ESCODEDO, TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION:
SOPTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/:
FILING DATE: January 11, 11
CLASSIFICATION: 514
                                                                                                          ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584
FILING DATE: January 11, 1996
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                 STREET: 633 West FI
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                                                                           COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                                                                                                                                      CITY: Los Angeles
STATE: California
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nes 10; Conservat
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TOPOLOGY: lir
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                                                                                                                                                                        COUNTRY:
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                                                                                                                                                                                                                                      633 West Fifth Street
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                                                                                                                                                                                                                                                                                                                                                                                              McSwiggen, James
Stinchcomb, Dan T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/08584040
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                                                                                           IBM Compatible
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                January 11, 1996
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OF VASCULAR ENDOTHELIAL
GROWTH FACTOR
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                                                                              IBM P.C.
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                                                                                                                                                                                                                                                                                       8502
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                                  US/08/584,040
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                                                                              DOS 5.0
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Gaps

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RESULT 74
US-08-584-040-7674
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
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NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: St
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LENGTH: 17 base pairs
TYPE: nucleic acid
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 216
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                  NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
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                                                      TELEPHONE: (213)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/584,040 FILING DATE: January 11, 1996 CLASSIFICATION: 514
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (213) 955-0440
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2332 TEGTECTTEGGGGTGTT 2348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7674, Application US/08584040
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Similarity 47.1%;
B; Conservative
                                       67-3510
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633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 U.S.A.
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                                                                                                            (213)
                                                             955-0440
                                                                                                        489-1600
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Pred. No. 8
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                                                                                                                  US-08-584-040-7684
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US-08-584-040-7684
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                                                                                                                                                                                                                                                     SOFTMARE: Word Perfect 5.1
SOFTMARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIPICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 3218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                 Query Match
Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No.
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Best Local Similarity 76.5%;
Matches 13; Conservative
                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
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APPLICANT: Escobedo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Pavco, Pamela APPLICANT: McSwiggen, James
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 633 West Fifth Street
STREET: 630 West Fifth Street
STREET: Suite 4700
                                                                                                                                            LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM COMPONERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: Suite 4700
CITY: Los Angeles
                                                                                                                                                                                                                               TELEFAX: (213) 955-0440 TELEX: 67-3510
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                                                                                                                                  TOPOLOGY:
2353 TGGGAGATCTTCACTTT 2369
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                                                  Conservative
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                                                                                                                              linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM Compatible
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                                                                1.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IBM P.C. DOS 5.0
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                                                 Score 13.8; D
Pred. No. 87;
6; Mismatches
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Pred. No. 87;
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                                                                                 DB 1;
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                                                  2
                                                                               Length 17;
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RESULT 77
US-08-584-040-7686
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                                                                                                                                                               Sequence 7686, Application US/08584040 Patent No. 6346398
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                                    GENERAL INFOR
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DO
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/58
PILING DATE: January 11, 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Stinchom
APPLICANT: Escobedo,
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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TELEX: 67-3510
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TELEFAX: (213) 955-0440
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                        2355 GGAGATCTTCACTTTAG 2371
                                                                                                   INFORMATION:
CANT: Pavco, Pamela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10;
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: Los Angeles
E: California
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Similarity 58.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   H: 17 base pairs nucleic acid
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633 West Fifth Street
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Stinchcomb, Dan T.
Escobedo, Jaime
McSwiggen, James
Stinchcomb, Dan T.
Escobedo, Jaime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/08584040
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Pred. No. 87
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US-08-584-040-7687
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Best Local Similarity 58.8%;
Matches 10; Conservative
                                                                                                                                                                                                                                                                                              sequence
                                                                                                                                                                                                                                                                              Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (213) 955-04
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                             TITLE OF INVENTION: N
TITLE OF INVENTION: TITLE OF INVENTION: (TITLE OF INVENTION: OTITLE OF INVENTION: OTITLE OF SEQUENCES:
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APPLICANT:
APPLICANT:
APPLICANT:
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LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/005,974 FILING DATE: October 26, 1995 ATTORNEY/AGENT INFORMATION:
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                                                                              CORRESPONDENCE ADDRESS:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS
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                              STREET:
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STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/584,040 FILING DATE: January 11, 1996 CLASSIFICATION: 514
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STATE: California
                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                  2356 GAGATCTTCACTTTAGG 2372
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10. 6346398
                                                                                                                                                                                                                                                           INFORMATION:
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California
                              Suite 4700
                                               3: Lyon & Lyon
633 West Fifth Street
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Escobedo, Jaime
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                                                                                                                                                                                                                                                                                              Application US/08584040
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                                                                                                                                                                           METHOD AND REAGENT FOR THE
                                                                                                                            TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL GROWTH FACTOR
                                                                                                                GROWTH FACTOR
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Pred. No. 87;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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COUNTRY: U.S.A.

90071-2066

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US-09-370-644B-21
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RESULT 80
US-09-474-432B-385/c
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LENGTH: 17
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                                                                                                                                                 Matches
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Best Local Similarity 58.8%;
                                                                                                                                                                                   Query Match
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/370,644B
CURRENT FILING DATE: 1999-08-06
PRIOR APPLICATION NUMBER: 08/596,257
PRIOR FILING DATE: 1996-04-18
NUMBER OF SEQ ID NOS: 26
SOFTMARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: KOSSMANN et al.

TITLE OF INVENTION: DEBRANCHING ENZYMES AND DNA SEQUENCES CODING THEM,
TITLE OF INVENTION: SUITABLE FOR CHANGING THE DEGREE OF BRANCHING OF
TITLE OF INVENTION: AMYLOPECTIN STARCH IN PLANTS
FILE REFERENCE: 514413-3771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENERAL INFORMATION:
                                                                                                                                                                                                                                     TYPE: RNA
ORGANISM: Solanum tuberosum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: January 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,97/
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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OPERATING SYSTEM:
                                                                                                                                                              Match 1.0%; Local Similarity 70.6%;
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                                                                                                            2271 AGTCAAGTGGATGGCTC 2287
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                                                                          AUUCAAGUGGAUGGCGC 17
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                                                                                                                                                                Score 13.8;
Pred. No. 8
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Pred. No. 87
                                                                                                                                                 Mismatches
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; SEQ ID NO 778
; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-474-432B-778
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US-09-474-432B-778
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SOFTWARE: PatentIn version
SEQ ID NO 385
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                                                                                                                                                                                                      APPLICANT: Sweedler, David
APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleotide triphosphate and their incorporation into oligonucleot
FILE REFERENCE: MBHB00-831-B (247/276)
CURRENT APPLICATION NUMBER: US/09/474,432B
CURRENT FILING DATE: 1999-12-19
PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR FILING DATE: 1997-11-05
PRIOR APPLICATION NUMBER: US 60/084,727
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR FILING DATE: 1998-11-04
PRIOR FILING DATE: 1998-11-04
PRIOR FILING DATE: 1998-11-04
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TITLE OF INVENTION: Nuclectide triphosphate and their incorporation into oligonuclect
FILE REFERENCE: MBHB00-831-B (247/276)
CURRENT APPLICATION NUMBER: US/09/474,432B
CURRENT FILING DATE: 1999-12-19
                                                                                                                 NUMBER OF SEQ ID NOS: 1526
SOFTWARE: PatentIn version
                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/301,511 PRIOR FILING DATE: 1999-04-28
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PRIOR FILING DATE: 1997-11-05
PRIOR APPLICATION NUMBER: US 60/084,727
PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR FILING DATE: 1998-11-04
PRIOR FILING DATE: 1998-11-04
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APPLICANT: Beigelman, Leo
APPLICANT: Burgin, Alex
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Adamic, Jasenka
Sweedler, David
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Burgin, Alex
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Adamic, Jasenka
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Beaudry, Amber
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                                                                                                                      version 3.0
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Pred. No. 8
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US-09-371-772B-1973
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US-09-371-772B-1973
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                                                                              PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1973
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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Best Local Similarity 58.8%;
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Patent No. 6566127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1973, Ap
Patent No. 656612
                                                                                                                                                                                                                              APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
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APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MSHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
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PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
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SOFTWARE: PatentIn version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
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APPLICANT: Pavco, Pam
               TYPE: RNA
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
                                                            ENGTH: 17
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Pred. No. 87
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Pred. No. 87;
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Query Match

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Score 13.8;

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; TYPE: RNA; Homo sapiens; ORGANISM: Homo sapiens
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US-09-371-772B-2124
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US-09-371-772B-2009
                                                                                                                                        SEQ ID NO 2124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 6566127
GENERAL INFORMATION:
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Query Match
Best Local Similarity
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                                                                                                                                                                     APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of I
TITLE OF INVENTION: Method and Reagent for the Treatment of I
TITLE OF INVENTION: Method and Reagent for the Treatment of I
TITLE OF INVENTION: Levels of Vascular Endothelial Growth F;
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT FILING DATE: 1996-US-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER: OF SEQ ID NOS: 14225
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CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
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APPLICANT: Pavco, Pam
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ORGANISM: Homo sapiens
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                                                                                                                   ENGTH:
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Stinchcomb, Dan
Escobedo, Jaime
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  Score 13.8;
Pred. No. 87;
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Pred. No. 8
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                  Length 17;
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APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
FITTLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
FITTLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/93/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR REPLING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PASTENTIAL Version 3.0
SEQ ID NO 2682
LENGTH: 17
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Query Match 1.0%;
Best Local Similarity 47.1%;
Matches 8; Conse, vative
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Best Local &
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                                                                                        TYPE: RNA
ORGANISM: Mus sp.
-09-371-772B-2682
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CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 60/05,974
PRIOR FILING DATE: 1995-10-26
PRIOR SPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
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TYPE: RNA
ORGANISM: Mus
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10; Conservative
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McSwiggen, Jim
Stinchcomb, Dan
Escobedo, Jaime
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Escobedo, Jaime
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      Score 13.8; I
Pred. No. 87;
7; Mismatches
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Pred. No. 8
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                                            DB 1; Length 17;
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; SEQ ID NO 3469
; LENGTH: 17
; LENGTH: 17
; TYPE: RNA
; ORGANISM: Mus sp.
US-09-371-772B-3469
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US-09-371-772B-3469
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US-09-371-772B-3459
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LENGTH: 17
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APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
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Matches 13; Conserv
                                            Query Match
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CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
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  Matches
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TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MedhBo0, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
                                                                                                                                                                                         PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
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APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
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Local Similarity les 9; Conserv
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52.9%;
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    6; Mismatches
                                         Score 13.8;
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                        Pred. No.
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                                            Length 17;
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                                                                                                                           ; TYPE: RNA
; ORGANISM: Mus sp.
US-09-371-772B-3471
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; ORGANISM: Mus sp.
US-09-371-772B-3470
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Best Local Similarity 58.8
Matches 10; Conservative
                                Query Match
Best Local Similarity
Matches 10; Conserv
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                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions |
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-10-8
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Patent No. 6566127
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PRIOR TILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
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APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371, 772B
CURRENT APPLICATION DATE: 1999-08-10
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                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 14225
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  2356 GAGATCTTCACTTTAGG 2372
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Escobedo, Jaime
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McSwiggen, Jim
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58.8%; Pred. No. 8
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                                                             Score 13.8;
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; TYPE: RNA
; ORGANISM: Mus sp.
US-09-371-772B-3472
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US-09-371-772B-4816
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                                                        Query Match
Best Local Similarity
Matches 13; Conserv
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SEQ ID NO 3472
LENGTH: 17
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APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions:
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-U (237/198)
CURRENT APPLICATION NUMBER: US/99/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 88/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
RIGHT STEAM OF SEQ ID NOS: 14225
NUMBER OF SEQ ID NOS: 14225
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TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-U (2377198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PILING DATE: 1995-10-26
PRIOR PILING DATE: 1995-01-08
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APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
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                                                                                                                                                               TYPE: RNA
ORGANISM: Homo sapiens
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2098 CAGCTGGCCAGAGGCAT 2114
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McSwiggen, Jim
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Escobedo, Jaime
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                                                                 Conservative
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                                                                                1.0%; Score 13.8; 76.5%; Pred. No. 8
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                                                                 Mismatches
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                                                                                      Query Match
Best Local Similarity
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                                                                       Matches
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APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-U (237/198)
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CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
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PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
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CURRENT FILING DATE: 1999-08-10
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SOFTWARE: PatentIn version 3.0
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TYPE: RNA
ORGANISM: Homo sapiens
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TYPE: RNA
ORGANISM: Homo sapiens
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Escobedo, Jaime
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                                                                                     76.5%;
                                                                  ; Score 13.8; D; Pred. No. 87; 2; Mismatches
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Pred. No. 87
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APPLICANT: Prescott
                              REFERENCE/DOCKET NUMBER: 20.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: NUMBER OF SEQUENCES:
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; TYPE: RNA
; ORGANISM: Human immunodeficiency virus
US-09-325-601-1
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PRIOR APPLICATION DATA: described beplication NUMBER: 08/141,244
FILING DATE: October 22, 1993
APPLICATION NUMBER: 07/953,681
FILING DATE: September 29, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Douglas E. Olson
REGISTRATION NUMBER: 22,798
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/325,601
CURRENT FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 53
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                                                                                                                                                                                                                     COMPUTER: IBM compatible
OPERATING SYSTEM: PC-DOS (Version 5
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/143,219
FILLING DATE: October 25, 1993
CLASSIFICATION: 435
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
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ADDRESSEE: Lyon & Lyon
                                                                                                                                                                          PRIOR APPLICATION DATA: PRIOR APPLICATION DATA:
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les 13; Conservative
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STREET: 611 West Sixth Street
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                                                                                                                                                        including application described below:
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Pred. No. 87;
2; Mismatches
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Query Match 1.0
Best Local Similarity 88.2
Matches, 15; Conservative

1.0%;

Score 13.8; Pred. No. 98; Mismatches

DВ 1;

Length 18

Indels

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1673 AACTTCCAGAGGACCCA 1689

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Best Local (
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APPLICANT: Winter, Grego
APPLICANT: Griffiths, Andrew
APPLICANT: Smith, Andrew
APPLICANT: Waterhouse, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                    TELEPHONE: 312-474-630 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
                                                                                                                                            FILING DATE: 15-MAY-1992
ATTORNEY/ACENT INFORMATION:
NAME: David W. Clough
REGISTRATION NUMBER: 36,107
REFERENCE/DOCKET NUMBER: 28
                                                                                                                                                                                                                                FILING DATE: 24-MAR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB92/00883
                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: GOIN PRIOR APPLICATION DATA:
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ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                         TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: PCT/GB93/00605
FILING DATE: 24-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/0:
FILING DATE: 16-SEP-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JMBER OF SEQUENCES:
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Local Similarity 88.2%;
nes 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TLE OF INVENTION: Methods for producing members of specific TLE OF INVENTION: binding pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INDIVIDUAL ISOLATE: COMPLEMENTARY TO THE RNA PROBE FOR INDIVIDUAL ISOLATE: PR-VII, FIGURE 5
                   STRANDEDNESS:
                                                 LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
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ropology:
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             nucleic acid
DEDNESS: single
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Smith, Andrew JH
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linear
                                                                                                      312-474-6300
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Best Local Similarity
Thes 15; Conserve
US-08-541-950B-13
             RESULT 100
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US-08-470-837-21/c
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                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 310-445-9031
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Sharp, Janice A.
REGISTRATION NUMBER: 34,051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE:
CLASSIFICATION: 424
                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: 310-445-1140
                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                            MOLECULE TYPE: cDNA
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                                                                                                                                                                                     NAME/KEY:
LOCATION:
                                                                                                                                                                                                                               NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 11150 Santa Monica Boulevard, Suite 400 CITY: Los Angeles
STATE: California
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ZIP: 90025-3395
                                                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 30630-1US01
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                                                                             1879 GAGATGATGAAGATGAT 1895
                                                       18 GTGATGATGATGAT 2
                                                                                                                                                                                                                                                                                                                     nucleic acid
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                                                                                                                                                                                                                                                                                                                                   18 base pairs
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                                                                                                                 Conservative
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                                                                                                               ο;
                                                                                                                             Score 13.8;
Pred. No. 9
                                                                                                               Mismatches
                                                                                                                                           DB 1;
                                                                                                                 2;
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                                                                                                                 Indels
                                                                                                               0
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Sequence 13, Application US/08541950B Patent No. 5821046

INFORMATION:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathle.
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 3777
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (617) 345-911 (NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM: MEDIUM TYPE: F,loppy disk
                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Johnson, Kevin Stuart
APPLICANT: Smith, Andrew John Hammond
TITLE OF INVENTION: Methods for producing members of specific
TITLE OF INVENTION: binding pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
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STATE: MA
CPAD/
                                                                                                                                                                                                               ADDRESSEE: David W. Clough
STREET: Marshall, O'Toole, Gerstein, Murray & Borun
STREET: 6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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13; Conserv
                                                                            60606-6402
                                                                                                                                                                              Chicago
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One Financial Center, 45th Floor
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                                                                                                                    USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nissim, Ahuva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Griffiths, Andrew David Williams, Samuel Cameron
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Waterhouse,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Winter, Gregory Paul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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SYSTEM: PC-DOS/MS-DOS
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76.5%; Pred. No. 98;
tive 2; Mismatches
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                                                    ; OTHER INFORMATION: Antisense Oligonucleotide US-09-205-921-17
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US-09-205-921-17/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-350-260A-79
                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: EX M. COWSETT
TITLE OF INVENTION: ANTISENSE MODULATION OF EGR-1 EXPRESSION
FILE REFERENCE: RTS-0028
FULE REPERENCE: RTS-0028
CURRENT APPLICATION NUMBER: US/09/205,921A
CURRENT FILING DATE: 1998-12-04
NUMBER OF SEQ ID NOS: 47
SEQ ID NO 17
                  Query Match
                                                                                                                                                                                                                                                                                                            Sequence 17, Application US/09205921A Patent No. 6008048
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                    ORGANISM: Artificial Sequence FEATURE:
                                                                                                                      TYPE: DNA
                                                                                                                                       ENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/307,619
FILING DATE: 16-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9110549.4
FILING DATE: 15-MAY-1991
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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FILING DATE: 15-MAY-
PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
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SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Clough, David W
REGISTRATION NUMBER: 36,107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
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SYSTEM: PC-DOS/MS-DOS
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15-MAY-1992
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88.2%;
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Pred. No. 98
Score 13.8; DB 1; Pred. No. 98;
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                Length 18;
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Best Local Similarity

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RESULT 104
US-09-344-521-25/c
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US-08-722-240-4
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SEQ ID NO 25
LENGTH: 18
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Patent No. 6100090
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                               APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowgert
TITLE OF INVENTION: ANTISENSE MODULATION OF PI3K P85 EXPRESSION
FILE REFERENCE: RTS-0062
FULR REFERENCE: RTS-0062
FULR REFERENCE: US/09/344,521
CURRENT APPLICATION NUMBER: US/09/344,521
CURRENT FILLING DATE: 1999-06-25
RUMBER OF SEQ ID NOS: 47
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk 3 1/2", 1.44 Mbyte COMPUTER: HP Vectra XU
OPERATING SYSTEM: Windows NT 4 Workstation SOPTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/722,240
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APPLICANT: van Mourik, Jan Aart
APPLICANT: Mertens, Koenraad
TITLE OF INVENTION: Method and of
TITLE OF INVENTION: disorders in
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REFERENCE/FOCKET NUMBER: St
TELECOMMUNICATION INFORMATION:
TELEPHONE: (732)530-6671
TELEFAX: (732)530-6584
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ATTORNEY/AGENT INFORMATION:
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328 Newman Springs Road,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Voorberg, Johannes Jacobus,
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                                                                                                                                                                                                                                                                                                                                                                     88.2%;
                                                                                                                                                                                                                                                                                                                                                   Pred. No. 98;
0; Mismatches
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, OTHER INFORMATION: Antisense Oligonucleotide US-09-344-521-25
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; OTHER INFORMATION: Antisense Oligonucleotide
US-09-205-143-41
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US-09-205-143-41/c
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LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
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Matches 15; Conservative
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APPLICANT: Lex M. Cowsert
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Matches 15; Conserv
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CURRENT FILING DATE: 1998-12-03
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                 ATTORNEY/AGENT INFORMATION:
NAME: WILLIAMS, Ph.D., Kath.
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 37
                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WOO'dPerfect 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
STREET: One Financial Center, 45th Floor
                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
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STATE: Ma
Th. 02111
TRAD/
                                                                                      APPLICATION NUMBER: FILING DATE:
                                                                                                                                     APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                               CITY: Boston
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                                                                                                     08/541,950
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                                                     Kathleen M.
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                   3777/57347
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Pred. No. 9
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Pred. No. 98
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                     (MRC-011AX)
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                                          REGISTRATION NUMBER: 36,107
REFERENCE/DOCKET NUMBER: 2811
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-474-6300
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity Matches 13; Conserv
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INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 60606-6402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                             APPLICATION NUMBER: PCT/GB92/00883
FILING DATE: 15-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: David W. Clough
                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO) CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/050,783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: synthetic RNA
                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 24-MAR
                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: 24-MAF
                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: 16-SE
               STRANDEDNESS:
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TOPOLOGY:
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linear
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fower, 233
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                                                                                                                                28111/32238
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South Wacker Drive
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Query Match
Best Local Similarity
Matches 15; Conserve
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US-09-104-337A-79
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; NAME/KEY: CDS
; LOCATION: (1)...(18)
US-08-868-452-21
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APPLICANT: Marcel E. Nimni
APPLICANT: Frederick L. Hall
APPLICANT: Lingtao Wu
                                                                                                                                                                                                                                                                                                                                                                                              Sequence 79, Applicati Patent No. 6492160 GENERAL INFORMATION:
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LENGTH: 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE; 17972-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: BONE MORPHOGENETIC PROTEINS AND THEIR TITLE OF INVENTION: USE IN BONE GROWTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Bo Han
APPLICANT: Edwin Shors
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Audrey L. Bartnicki
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                           Johnson, Kevin Stuart
Smith, Andrew John Hammond
TITLE OF INVENTION: Methods for producing members of specific binding pairs
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                                                                                                                                                               STREET: Marshall, C
                                                                                                                               STATE: Illinois
                                                                                                                                                CITY: Chicago
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                                                                                                                COUNTRY: USA
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                                                                                                60606-6402
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hs, Andrew David
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s Tower, 233 Sout!
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Pred. No. 98
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Pred. No. 9
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                                                                                                                                                                 South Wacker Drive
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APPLICATION NUMBER: US/09/104,337A

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; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Designated
; OTHER INFORMATION: a forward oligonucleotide encoding (His)6
US-09-280-030-8
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Best Local Similarity
                                             Query Match
                                                                                                                                                                                                                                  SEQ ID NO 8
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      Matches
                                                                                                                                                                                                                                                                                                                               APPLICANT: Sato, Seiji
APPLICANT: Higashikuni, Naohiko
APPLICANT: Kudo, Toshiyuki
APPLICANT: Kondo, Masaki
APPLICANT: Kondo, Masaki
APPLICANT: Kondo, Masaki
TITLE OF INVENTION: DNAS ENCODING NEW FUSION PROTEINS AND PROCESSES FOR
TITLE OF INVENTION: DNAS
TITLE OF INVENTION: DNAS
FILE REFERENCE: 382.1026
CURRENT APPLICATION NUMBER: US/09/280,030A
CURRENT FILING DATE: 1999-03-26
                                                                                                                                                                                                                                                                         CURRENT FILING DATE: 1999-03-26
EARLIER APPLICATION NUMBER: JP10-87339/1998
EARLIER FILING DATE: 1998-03-31
NUMBER OF SEQ ID NOS: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENERAL INFORMATION:
                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                               LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 79:
    Local Similarity 88.2%; es 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Bartnicki, Audrey L.
REGISTRATION NUMBER: 40,499
REFERENCE/DOCKET NUMBER: 28111/32372A
TELECOMMUNICATION INFORMATION:
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TYPE: nucleic acid
STRANDEDNESS: single
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APPLICATION NUMBER: US 08/150,002
FILING DATE: 31-MAR-1994
APPLICATION NUMBER: US 08/307,619
FILING DATE: 16-SEP-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/350,260 FILING DATE: 05-DEC-1994
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APPLICATION NUMBER: GB 9206318.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 24-MAR-1992
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  Conservative
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Pred. No. 98
                       Score 13.8;
Pred. No. 98
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                                         Length 18;
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                                                                                                                                      ; SEQ ID NO 3
; LENGTH: 1B
; TYPE: RNA
; ORGANISM: Human immunodeficiency virus
US-09-325-601-3
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US-09-280-030-9
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                                                                Query Match
Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 3, Application US/09325601 Patent No. 6573045
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Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 9, Application US/09280030A Patent No. 6506595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/325,601 CURRENT FILING DATE: 1999-06-03
                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Methods and Kits for Discovery of RNA-Binding Compounds FILE REFERENCE: 3950/81235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Sato, Seiji
APPLICANT: Higashikuni, Nac
APPLICANT: Kudo, Toshiyuki
APPLICANT: Kondo, Masaaki
                                                                                                                                                                                                                                                                                                                                             APPLICANT: Karn
APPLICANT: Prescott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EARLIER APPLICATION NUMBER: JP10-87339/1998
EARLIER FILING DATE: 1998-03-31
NUMBER OF SEQ. ID NOS: 66
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 382.1026
CURRENT APPLICATION NUMBER: US/09/280,030A
CURRENT FILING DATE: 1999-03-26
                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: DNAS ENCODING NEW FUSION PROTEINS AND PROCESSES FOR TITLE OF INVENTION: PREPARING USEFUL POLYPEPTIDES THROUGH EXPRESSION OF TITLE OF INVENTION: DNAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: Designated OTHER INFORMATION: a reverse oligonucleotide encoding (His)6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH: 18
                              1490 AGCCAGACTTCAGCAGC
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                                                                1.0%;
ilarity 76.5%;
Conservative
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                                                                                     Score 13.8;
Pred. No. 9
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Pred. No. 98;
                                                                    Mismatches
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RESULT 113 US-09-081-646-233/c ; Sequence 233, Application US/09081646

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; LENGTH: 15
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-081-646-233
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APPLICANT: Zhang, Lin
APPLICANT: Zhou, Wei
TITLE OF INVENTION: Gene Expression Profiles in
TITLE OF INVENTION: Cancer Cells
FILE REFERENCE: 01107.74664
CURRENT APPLICATION NUMBER: US/09/081,646
CURRENT FILING DATE: 1996-05-20
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ ID NOS: 871
NUMBER OF SEQ ID NOS: 871
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                                                       FILING DATE: NO. 5612215e
APPLICATION NUMBER: 07/98
FILING DATE: December 7,
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                   OPERATING SYSTEM: IBM P.C. DOS SOFTWARE: FASTSEQ Version 1.5 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
NAME: Warburg, Richard REGISTRATION NUMBER: 32 REFERENCE/DOCKET NUMBER:
                                                                                                                                   FILING DATE: December 13, 1994 APPLICATION NUMBER: 08/152,487
                                                                                                                                                                                                           APPLICATION NUMBER: US/08/390,850 FILING DATE: February 17, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 1.0%;
Local Similarity 93.3%;
                                                                                                                                                                           APPLICATION NUMBER: 08/354,920
                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
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633 West Fifth Street
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VENTION: METHOD AND REAGENT FOR TREATMENT
VENTION: OF ARTHRITIC CONDITIONS
                                                                                                                                                                                                                                                                                                                                                                                                        U.S.A.
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Gustofson, John
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                                                                                                                                                                                                                                                                                                             IBM Compatible
                                                                                                                 No. 5612215ember 12,
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                                                                                                 07/989,848
                 32,327
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HYPOTHETICAL:
; ANTI-SENSE: 1
US-08-398-008A-5
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US-08-398-008A-5/c
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Best Local S
Matches 12
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                                                            Matches
                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/126,501
APPLICATION NUMBER: 08/126,501
FILING DATE: 24-SEP-193
AITORNEY/AGENT INFORMATION:
NAME: Adler, Dr. Benjamin Aaron
REGISTRATION NUMBER: 35,423
REFERENCE/DOCKET NUMBER: 05705CIP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (713) 777-2321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: MACINTOSH IICI
OPERATING SYSTEM: Macintosh
SOFTWARE: Microsoft Word 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/398,
FILING DATE: March 2, 1995
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MEDIUM TYPE: Diskette, 3
COMPUTER: MACINTOSH IIC1
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                                                                                                                                                    MOLECULE TYPE: DNA HYPOTHETICAL: no
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nes 12; Conserv
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STRANDEDNESS: single
                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                 STRANDEDNESS:
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16 GTGATTACAGCTTCC
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                                                            Conservative
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                                                                            Score 13.4; DB 1;
Pred. No. 1e+02;
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Pred. No. 1
                                                            Mismatches
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                 Sequence 5, Application US/08893333A Patent No. 5981705
                                                                                                                                                                                                                                  Query Match
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 514
PRIOR APPLICATION UNBER: 08/390,850
FILING DATE: Pebruary 17, 1995
APPLICATION NUMBER: 08/354,920
FILING DATE: December 13, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 211/084
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: December 7,
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:
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MEDIUM TYPE: 8torage
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CORRESPONDENCE ADDRESS:
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APPLICANT: Gustofson, John
TITLE OF INVENTION: METHOD AND REAGENT FOR TREATMENT
TITLE OF INVENTION: OF ARTHRITIC CONDITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                  Local Similarity
les 12; Conserv
                                                                                                                                                                                                                                                                                             TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/152,487 FILING DATE: No. 5731295ember 12, APPLICATION NUMBER: 07/989,848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/01
FILING DATE: 05-MAY-1995
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                                                                                                                                                             1586 TGAĄCTCCĄĄCĄCCC 1600
INFORMATION:
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California
                                                                                                                         UGGACUCCAÁCACCC 17
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                                                                                                                                                                                              Score 13.4; DB 1; Length 17; Pred. No. 1e+02; 2; Mismatches 1; Indels
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SEQ ID NO 5
                                                                                                                                                                                                                                    COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/776,900C
FILING DATE: 30-APR-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: PCT/FR95/01035
FRIOR APPLICATION NUMBER: PCT/FR95/01035
FILING DATE: 01-AUG-1995
PRIOR APPLICATION NUMBER: FR/94/09567
FILING DATE: 02-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: CHARLES A. MUSERLIAN
REGISTRATTON NUMBER: 10-683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 10, Application US/08776900C Patent No. 6020477
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: DNA Encoding Natural Killer Lytic Associated Protein FILE REFERENCE: D5705CIP/D
                                                                               TELEFAX: (212) 661-800
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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TYPE: DNA
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OTHER INFORMATION: Used to sequence Natural Killer Lytic Associated Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: primer_bind
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: DIU, Anita; FAUCHEU, Chi; Hercend, Thierry; APPLICANT: LALANNE, Jean-Louis, LIVINGSTON, David and APPLICANT: SU, Michael
TITLE OF INVENTION: DNA SEQUENCES CODING FOR THE HUMAN TITLE OF INVENTION: PROTEINS TX AND TY RELATED TO THE TITLE OF INVENTION: INTERLEUKIN-1BETA COMPREDENCE.
                                                                                                                                                            NAME: CHARLES A. MUSERLIAN REGISTRATION NUMBER: 19,683 REFERENCE/DOCKET NUMBER: 146.1265 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: BIERMAN & MUSERLIAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
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STATE: NEW YORK
COUNTRY: USA
TOPOLOGY:
                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 10016
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                                            nucleic acid
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                                                                                                                                  (212) 661-8002
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                                                                                                                                                       (212) 661-8000
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30-APR-1997
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                  single
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                                                                                                                                                                                                                                                                                                                                                                                  PCT/FR95/01035
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SEQ ID NO: 1 from 695 to

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RESULT 120
US-08-584-040-4355
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Matches
                                                                                          Patent No.
                                                                                            Sequence 4355, Application US/08584040 Patent No. 6346398
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                                                            GENERAL INFORMATION: APPLICANT: Pavco,
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               APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: lin
MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (OEB)
SOFTWARE: + Corrections under WORDPERFECT 5.1 for SEQ ID NO 22
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
IPPLICANT: Escobedo, Jaime
                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
LOCATION: 1..17
OTHER INFORMATION: /note= "SEQ ID NO 1 FROM 695 TO 711"
                                                                                                                                                                                                                                                              Match 1.0%;
Local Similarity 93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: ROUSSEL UCLAF
STREET: 102, Route de No. 61803868y
CITY: ROMAINVILLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 15-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
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                                                                                                                                                                                                                 2465 AACTGTACATGATGA 2479
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                                                                                                                                                                                                                                                14; Conservative
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VENTION: DNA SEQUENCES CODING FOR THE HUMAN
SEQUENCES: 42
              McSwiggen, James
Stinchcomb, Dan T.
Escobedo, Jaime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FRANCE
                                                            Pavco, Pamela
                                                                                                                                                                                                                                                                                                                                                                                     Other nucleic acid /desc = "OLIGONUCLEOTIDE"
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DMBER: 776,900
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Pred. No. 1e+02;
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Pred. No. 16
                                                                                                                                                                                                                                            Mismatches
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le+02;
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                                                                                                                                                                                                                                                                                         Sequence 4356, App. Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
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APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stonchoomb, Dan T.
APPLICANT: Stonchoomb, Dan T.
APPLICANT: Stonchoomb, Dan T.
APPLICANT: Stinchoomb, Dan T.
APPLICANT: Stinchoomb, Dan T.
APPLICANT: Stinchoomb, Dan T.
APPLICANT: Stinchoomb, James
TITLE OF INVENTION: GROWTH FACT
TITLE OF INVENTION: GROWTH FACT
                                                                                                                                                                                                                                                                             GENERAL
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                    NERAL INFORMATION:
APPLICANT: Pavco, Pamela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 514
PRIOR APPLICATION NUMBER: (
                                                                                     CORRESPONDENCE ADDRESS
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CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 1211
TELEFAX: 67-3510
TELEX: 67-3510
                              STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: Suite 470
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
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STREET: 633 West Fifth Street
                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
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Los Angeles
: California
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                                 Suite 4700
                                                   633 West Fifth
                                                                                                                                                                                                                                                                                                         Application US/08584040
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         January 11, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.0%;
                                                                                                                                                      METHOD AND REAGENT FOR THE TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS
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                                                                                                                    GROWTH FACTOR
                                                                                                                                      OF VASCULAR ENDOTHELIAL
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                                                   Street
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US-09-474-432B-649
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                                                                                                                     PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR FILING DATE: 1997-11-05
PRIOR APPLICATION NUMBER: US 60/084,727
PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR APPLICATION NUMBER: US 09/301,511
PRIOR FILING DATE: 1998-11-04
PRIOR FILING DATE: 1999-04-28
NUMBER OF SEQ ID NOS: 1526
                                                                                  SOFTWARE: PatentIn version 3.0 SEQ ID NO 649 .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                    APPLICANT: Sweedler, David
APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleotide triphosphate and their incorporation into oligonucleot
FILE REFERENCE: MBHB00-831-B (247/276)
CURRENT APPLICATION NUMBER: US/09/474,432B
CURRENT FILING DATE: 1999-12-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: October 26, 1995
ATTOREY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Beigelman, Leo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (213) 955-04
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                 TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid STRANDEDNESS: sing TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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Beaudry, Amber
Karpeisky, Alex
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Adamic, Jasenka
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.0%; Score 13.4; DB 1; Length 17; 60.0%; Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM P.C. DOS 5.0
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Query Match

Score 13.4;

DB 1;

Length 17;

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                                                                                                                                                                                   TITLE OF INVENTION: Method and Reagent for the Treatme:
TITLE OF INVENTION: Levels of Vascular Endothelial Gr.
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-08
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US-09-371-772B-2123
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; ORGANISM: Homo sapiens
US-09-371-772B-2122
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US-09-371-772B-2122
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LENGTH: 17
Query Match
Best Local Similarity
                                                                                                                                            SEQ ID NO 2123
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Patent No. 6566127
GENERAL INFORMATION:
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GENERAL INFORMATION:
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Best Local :
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR PPLICATION NUMBER: US 08/584,040
PRIOR PILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Pavcó, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
                                                                                                                                                                 SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.0
                                                                           LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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Local Similarity 60.0%;
les 9; Conservative
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Stinchcomb, Da
                                                                                                                                                                                                                                                                                                                                                                                                  Escobedo, Jaime
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1.0%;
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Pred. No. 1e+0
5; Mismatches
Score 13.4;
Pred. No. 1e
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1e+02;
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RESULT 126
5185440-12
;Patent No. 5185440
; APPLICANT: DAVIS, NANCY L.;WILLIS, LORETTA V.;JOHNSTON,
;ROBERT E.;SMITH, JONATHAN F.
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                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (212) 751-68' INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "synthetic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version SOFTWARE: #1.30 (EPO) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 35,353
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 311130/94
FILING DATE: 21-NOV-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/765,34
FILING DATE: 23-DEC-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 145146/94
FILING DATE: 27-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: MATSUDA, Y.,
APPLICANT: KONDO, S.
TITLE OF INVENTION: AN ANTISENSE NUCLEIC ACID
TITLE OF INVENTION: COMPOUND
                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (212)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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NEW YORK
                                                                                                                                                                                                                                                                                                                                                nucleic acid
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UCHIDA, T.,
TANAKA, Y.,
MATSUDA, Y.,
KONDO °
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
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                                                                                                                                                                                                                                                                                                                                                                                                                 (212) 758-4800
12) 751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Floppy disk
                                                                                                                                                                                                                                                                                                                                single
                                                                                                                                                                                          0.9%; Score 13; DB 1; Length 14; 100.0%; Pred. No. 80; tive 0; Mismatches 0; Indels
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   Query Match
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US-08-390-850-566
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: CDNA CLONE CODING FOR VENEZUELAN EQUINE ENCEPHALITIS VIRUS AND ATTENUATING MUTATIONS THEREOF NUMBER OF SEQUENCES: 14
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO:12:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/390,850
FILING DATE: February 17, 1995
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/354,920
FILING DATE: December 13, 1994
APPLICATION NUMBER: 08/152,487
FILING DATE: No. 5612215ember 12, 1993
APPLICATION NUMBER: 07/989,848
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Stinchcomb, Dan T.
TITLE OF INVENTION: METHOD AND REAGENT FOR TREATMENT
TITLE OF INVENTION: OF ARTHRITIC CONDITIONS
                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
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FILING DATE: 20-JUN-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. DO: SOFTWARE: FastSEQ Version 1.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 14
                                                                                                                                                                                                    NAME: Warburg, Richard REGISTRATION NUMBER: 32, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                   TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 90071
                                                                      LENGTH:
                                                                                                                                               TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: Suite 4700
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                                                   nucleic acid
                                                                                                                             67-3510
                                                                      17 base pairs
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                                                                                                                                                (213) 955-0440
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Gustofson, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Draper,
               linear
                                  single
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                                                                                                                                                                                                                 32,327
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80;
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0.9%;

Score 13;

DB 1;

Length 17;

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RESULT 129
US-08-435-634-566/c
; Sequence 566, Application US/08435634
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US-08-390-850-567/c
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                                                                                                                                                                                                                                    US-08-390-850-567
                                                                                                                                                           Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                              TELEX: 67-3510
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: February 17, 1995
PRICATION DATA:
APPLICATION NUMBER: 08/354,920
FILING DATE: December 13, 1994
APPLICATION NUMBER: 08/152,487
FILING DATE: No. 5612215ember 12, 1:
APPLICATION NUMBER: 07/989,848
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Stinchcomb, Dan T.
TITLE OF INVENTION: METHOD AND REAGENT FOR TREATMENT
TITLE OF INVENTION: OF ARTHRITIC CONDITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                        TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Warburg, Richard REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
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                                                                                                                          2416 CTGCTGAAGGAAG 2428
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Gustofson, John
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                                                                   APPLICANT:
                                                      APPLICANT:
OF INVENTION: METHOD AND REAGENT FOR TREATMENT
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RESULT 130
US-08-435-634-567/c
                                                  Sequence 567, Application US/08435634 Patent No. 5731295
                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                     GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
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APPLICANT:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/390,850
FILING DATE: Pebruary 17, 1995
APPLICATION NUMBER: 08/354,920
FILING DATE: December 13, 1994
APPLICATION NUMBER: 08/152,487
FILING DATE: No. 5731295ember 12,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/435,634 FILING DATE: 05-MAY-1995 CLASSIFICATION: 514
                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Warburg, Richard
REGISTRATION NUMBER: 3
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STREET: 633 West Fifth Street
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California
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Draper, Kenneth G.
Pavco, Pamela
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Gustofson, John
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100.0%; Pred. No. 1.2e+02;
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McSwiggen, James Gustofson, John

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RESULT 131
US-08-479-614-18/c
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Best Local Similarity 100.0%; P
Matches 13; Conservative 0;
                                                                                                                                                                                                                                              Sequence 18, Application US/08479614
Patent No. 5861294
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
COMPUTER: Macintosh
                                                                                                                                                          NUMBER OF SEQUENCES: 3
                                                                                                                                                                                           APPLICANT: Cowart, Marlon Daniel, Halbert, Donald N., APPLICANT: Kerwin, Jr., James F., McNally, Teresa TITLE OF INVENTION: Adenosine Kinase Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Lon nor 4
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APPLICATION NUMBER: 08/354,920
FILING DATE: December 13, 1994
APPLICATION NUMBER: 08/152,487
FILING DATE: No. 5731295ember 12,
APPLICATION NUMBER: 07/989,848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Warburg, Richard REGISTRATION NUMBER: 32,327 REFERENCE/DOCKET NUMBER: 211/084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 05-MAY
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                                                                                                      CITY: Abbott Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: December
                                                                     COUNTRY:
                                                                                                                         STREET:
                                                                                                                                           ADDRESSEE:
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                                                                                                                     E: Abbott Laboratories
D-377 AP6D, 100 Abbott Park Road
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SYSTEM: IBM P.C. DOS 5.0
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ry 17, 1995
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                  3.50 inch
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                                                         ZIP: 90071-2066

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Thomas D. Brainard
REGISTON NUMBER: 32,459
REFERENCE/DOCKET NUMBER: 57.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (708) 937-4884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 536
PRIOR APPLICATION DATA:
            CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/001,135
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Folkerts, Otto
APPLICANT: Merlo, Donald J.
TITLE OF INVENTION: COMPOSI
TITLE OF INVENTION: MODULAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Microsoft Word 6.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                           STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   . 2400 GGAGGAACTTTTAA 2414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                         COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: 811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: June 7, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16 GRAGRAAYTTYTTRA 2
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Young, Scott A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Edington, Brent E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                              Lyon & Lyon
July 13, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patricia Ann Owens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Thomas A.
                                                                                                                                                                                                                                                                                                                                                                                  COMPOSITION AND METHODS FOR MODULATION OF GENE EXPRESSION IN PLANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%;
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Pred. No. 1.2e+0
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RESULT 134
US-09-371-772B-5850/c
Sequence 5850, Application US/09371772B
Patent No. 6566127
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                                                                                                                                                                                                                                                                                                                                                                                 ; LENGTH: 16
; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-371-772B-5819
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US-09-371-7728-5819
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TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 748:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 5819
                                                                                                                                                                                                                                                                                                       Matches 11;
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                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Scinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00, 876-J (237/198)
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
                    APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam APPLICANT: McSwiggen, Jim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
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TOPOLOGY: linear
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 0.9%;
Local Similarity 76.9%;
                                                                                                                                                                                                                                                   1820 TGAAGATGTTGAAAGA 1835
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McSwiggen, Jim
Stinchcomb, Dan
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Pred. No. 1.2e+02;
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Pred. No. 1.1e+02;
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US-08-179-738-23/c
                          RESULT 136
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; ORGANISM: Homo sapiens
US-09-371-772B-5850
                                                                                                                                                                                                                                                                                               SOPTWARE: PatentIn version 3.0
SEQ ID NO 7080
LENGTH: 16
TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                         Query Match
Best Local 8
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GENERAL INFORMATION:
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE REFERENCE: MSHB00,876-J (237/198)
CURRENT FALLING NOWBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PILING DATE: 1995-10-26
PRIOR PILING DATE: 1995-01-08
NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER: OF SEO ID NOS: 14225
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CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR TILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 60/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-10-26
PRIOR PILING DATE: 1995-10-80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
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les 14; Conserv
                                                                                                                                        1820 TGAAGATGTTGAAAGA 1835
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                                                                                           1 UCAAAAUGUUGAAAGA 16
                                                                                                                                                                                   l Similarity
10; Conserv
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McSwiggen, Jim
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Pred. No. 1.1e
0; Mismatches
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1.1e+02;
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Sequence 23, Application US/08179738
Patent No. 5578462
GENERAL INFORMATION:

APPLICANT: Seizinger, Bernd R.
APPLICANT: Kley, Nikolai A.
APPLICANT: Bianchi, Albert B.
TITLE OF INVENTION: No. 5578462el NF2 Isoforms
NUMBER OF SEQUENCES: 26

CORRESPONDENCE ADDRESS:

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US-08-390-850-694
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARR: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/179,738
FILING DATE: 10-JAN-1994
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Robins, Roberta L.
REGISTRATION NUMBER: 33,208
REFERENCE/DOCKET NUMBER: 3998-0017
TELECOMMUNICATION INFORMATION:
TELECHNUE: (415) 617-8999
TELECHNUE: (415) 617-8999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 87.9
Matches 14; Conservative
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APPLICATION NUMBER: US/08/390,850
FILING DATE: February 17, 1995
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/354,920
FILING DATE: December 13, 1994
                                                                                              COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P. C. DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: McSwiggen, James
APPLICANT: Gustofson, John
APPLICANT: Stinchcomb, Dan T.
TITLE OF INVENTION: METHOD AND REAGENT FOR TREATMENT
TITLE OF INVENTION: OF ARTHRITIC CONDITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: cDNA
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LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                              COMPUTER: IBM Com
OPERATING SYSTEM:
SOFTWARE: FastSEC
                                                                                                                                                                                                                                                                             CITY: Los Angeles
STATE: California
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TOPOLOGY: lir
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                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                        STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
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633 West Fifth Street
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) 327-3231
NO: 23:
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US-08-390-850-695
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Hest Local Similarity 75.0%;
Matches 12; Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 695, Application US/08390850
                                                                                          COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FASTESQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,850
FILING DATE: February 17, 1995
PRIOR APPLICATION NUMBER: 08/154,920
APPLICATION NUMBER: 08/154,920
APPLICATION NUMBER: 08/152,487
TILING DATE: December 13, 1994
APPLICATION NUMBER: 08/152,487
FILING DATE: No: 5612215ember 12, 19
APPLICATION NUMBER: 07/989,848
FILING DATE: December 7, 1992
ATTORNEY/ACENT INFORMATION:
NUMBER: 07/989,848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Draper, Kennet
APPLICANT: Pavco, Pamela
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: No. 5612215ember 12,
APPLICATION NUMBER: 07/989,848
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Stinchcomb, Dan T.
TITLE OF INVENTION: METHOD AND REAGENT FOR TREATMENT
TITLE OF INVENTION: OF ARTHRITIC CONDITIONS.
NUMBER OF SEQUENCES: 1151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 633 West STREET: Suite 470 CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Warburg, Richard REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                           90071
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633 West Fifth Street
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Gustofson, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss: single
linear
                                                                                                                                                                                                                                                                                                                                                         : storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      32,327
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                                                                                                                                                                                                                                                                                                                                              DOS 5.0
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Best Local Similarity
Marches 12; Conserve
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Patent No. 5646...

PATERAL INFORMATION:
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                                                                 TELEFAX: (213) 955-044
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/373,1
FILING DATE: January 13, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: McSwiggen, James
APPLICANT: JAYVIS, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
                                                                                                                                                                                                                      FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
FILING DATE: August 26, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                     TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                     TELEPHONE: (213) 955-0440
                                                                                                                                                                      NAME: Warburg, Richard REGISTRATION NUMBER: 32,327 REFERENCE/DOCKET NUMBER: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Con
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP:
          STRANDEDNESS:
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STATE: California
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                                                LENGTH:
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                            nucleic acid
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                                              17 base pairs
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633 West Fifth Street
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linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IBM P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                  US/08/373,124A
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US-08-373-124A-534
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                                                                                        Query Match
Best Local
                                                                     Matches
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Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/373,124
FILING DATE: January 13, 1995
PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
PILING DATE: August 26, 1992
APPLICATION NUMBER: 07/936,422
PILING DATE: August 26, 1992
ATTORNEY/ACENT INFORMATION:
NAME: Warburg, Richard
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: McSwiggen, James
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                TELEFAX: (41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                            0.9%;
Local Similarity 81.2%;
hes 13; Conservation
                                                                                                                                                    STRANDEDNESS: sir
                                                                                                                                                                                                                                                               TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
STATE: California
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                    1584 CATGAACTCCAACACC 1599
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CAAGAACUCCUACACC
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                                                                                    Score 12.8;
Pred. No. 1.3
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Pred. No. 1.3e+02;
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2;
                                                                                .3e+02;
                                                                                                     DB 1; Length 17;
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                                                                  Gaps
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RESULT 142
US-08-373-124A-1116/c
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                                                                                              Sequence 1116, Application US/08373124A Patent No. 5646042
                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/373,12
FILING DATE: January 13, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION UNMBER: 08/122,943
FILING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 08/12,943
FILING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 07/987,132
                  GENERAL INFORMATION:
APPLICANT: Stinch
APPLICANT: Draper
APPLICANT: McSwig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 :NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Jarvis, Thale
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 07/936,422 FILING DATE: August 26, 1992 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (213) 489-10
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
TOPOLOGY: li
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
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                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                       2240 ATTACAAAAAGACCAC 2255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION:
                                                                                                                                                                                                                      17 ATAACAAAAAAACCAC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Los Angeles
: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90071
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Draper, Kenneth
McSwiggen, James
Draper, Kenneth
McSwiggen, James
Jarwis, Thale
                                                         Stinchcomb, Dan T.
                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stinchcomb, Dan T.
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IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lyon & Lyon
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                                                                                                                                                                                                                                                                                                                                                                                                               single
                                                                                                                                                                                                                                                                                                                0.9%;
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                                                                                                                                                                                                                                                                                              0,
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                                                                                                                                                                                                                                                                                                                   Score 12.8; DB 1; Length 17; Pred. No. 1.3e+02;
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                                                                                                                                                                                                                                          Patent No. 5646042
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                       Sequence 2423, Application US/08373124A
                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/373,12.
FILING DATE: January 13, 1995
PRIOR APPLICATION DATA:
APPLICATION UNUBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
APPLICATION NUMBER: 08/192,943
APPLICATION NUMBER: 07/987,132
FILING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
FILING DATE: December 7, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 07/936,4 FILING DATE: August 26, 1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION:
                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyo
                                                                                                         TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                    APPLICANT:
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                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 9007:
                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (213) 489-160
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH:
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Los Angeles
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17 base pairs
                     Suite 4700
                                  8: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                               Draper, Kenneth
McSwiggen, James
Jarvis, Thale
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                                                                                                                                                                                                                         Stinchcomb, Dan T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%;
                                                                                                         METHODS AND COMPOSITIONS FOR TREATMENT OF RESTENOSIS AND CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            METHODS AND COMPOSITIONS FOI
TREATMENT OF RESTENOSIS AND
CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07/936,422
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Pred. No. 1.3e+02;
0; Mismatches 2;
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage

COUNTRY:

California

U.S.A.

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US-08-435-634-694
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 694, Application US/08435634
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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APPLICANT: Draper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: February 7, 1994
ARPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
FILING DATE: August 26, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
APPLICANT:
                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                          APPLICANT: McSwiggen, James
APPLICANT: Gustofson, John
APPLICANT: Stinchcomb, Dan T.
TITLE OF INVENTION: METHOD AND REAGENT FOR TREATMENT
TITLE OF INVENTION: OF ARTHRITIC CONDITIONS
                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REPERENCE/DOCKET NUMBER: 209/035
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: January PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
                                                                                                                                  ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ Version 1.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/373,124A
FILING DATE: January 13, 1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1584 CATGAACTCCAACACC 1599
                                                                                                                  90071
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                                                                                                                                                                                                                                                                                                                                                                                              Pavco, Pamela
                                                                                                                                                                                                                                                                                                                                                                                                                Draper,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
                                                                                                                                                                                                                                                                                                                                                                                                                Kenneth G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%; Score 12.8; DB 1; Length 17; 81.2%; Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                         1151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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Best Local Similarity 75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/390,850
FILING DATE: February 17, 1995
APPLICATION NUMBER: 08/354,920
FILING DATE: December 13, 1994
APPLICATION NUMBER: 08/152,487
FILING DATE: No. 5731295ember 12, 19
APPLICATION NUMBER: 07/389,848
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Markhyr Bichard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (213) 955-044
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Pavco, Lumes
APPLICANT: McSwiggen, James
AppliCANT: Gustofson, John
 CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
FILING DATE: February
                                                                     COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS
SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,
FILING DATE: 05-MAY-1995
                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: METHOD AND REAGENT FOR TREATMENT TITLE OF INVENTION: OF ARTHRITIC CONDITIONS NUMBER OF SEQUENCES: 1151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 211/084
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                     CITY: Los Angeles
STATE: California
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                                                                                                                                                                                                                                 ZIP: 90071
                                                                                                                                                                                                                                                                                                      STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (213) 955-0440
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                695, Application US/08435634
o. 5731295
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                                                                                                                                                                                                                                                    U.S.A.
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February
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                  08/390,850
                                                                                       US/08/435,634
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                                                                                                                                             .C. DOS 5.0
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21, Application US/08623891 Patent No. 5795778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (213) 489-1
TELEFAX: (213) 955-044
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 211/084
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                   FILING DATE:
APPLICATION NUMBER: 07/882,921
FILING DATE: May 14, 1992
APPLICATION NUMBER: 07/948,359
FILING DATE: September 18, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: Wordperfect (Version 5.1)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 611 West Sixth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: December 7, ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 435
                                                 NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 200/209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                            APPLICATION NUMBER: US/07/987,133
                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 90017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Los Angeles
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UMBER: 07/989,848
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METHOD AND REAGENT FOR
INHIBITING HERPES SIMPLEX
VIRUS REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  er 13, 1994
08/152,487
                                                                                                                                                                                                                                                US/08/238,200
                                                                                                                                                                                                                                                                                                                      US/08/623,891
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US-08-758-306-123/c
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                                                      Matches
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Best Local Similarity
                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS
SOFTWARE: FASTSON DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/758,3
FILING DATE: December 3, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 212/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDENNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                        TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Stinchcomb, wan 1.
APPLICANT: McSwiggen, James A.
APPLICANT: McSwiggen, James A.
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: TREATMENT OF DISEASES
TITLE OF INVENTION: ASSOCIATED WITH
TITLE OF INVENTION: GAMMA-CHAIN EXPRESSION
TITLE OF INVENTION: GAMMA-CHAIN EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A. ZIP: 90071-2066
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2156 CAGCCAGAAATGTTTT 2171
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                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/08758306
                                                                                                                                                                                                                                                             (213) 955-0440
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linear
                                                      0.9%;
ilarity 87.5%;
Conservative
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75.0%;
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                                                                                                                                                                                                                                                                                                                212/132
                                                                      Score 12.8; DB 1;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 12.8; DB 1;
Pred. No. 1.3e+02;
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                                                          Mismatches
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TELEFAX: (212)
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
RESULT 149
US-08-435-628-534
; Sequence 534, Application US/08435628
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                                                                                                                                                                                                                                                                         US-08-435-628-416
                                                                                                                                                                                           Matches
                                                                                                                                                                                                       Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/435,620
FILING DATE: 05-MAY-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/373,124
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION WHERE: 07/936,422
FILING DATE: August 26, 1992
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Stinchcomb, Dan T.
APPLICANT: Draper, Kenneth
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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STATE: California
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                                                                                                                                        1866 GGTGTCAGAGATGGAG 1881
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633 West Fifth Street
                                                                                                                                                                                         Conservative
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                                                                                                                                                                                    Score 12.8; DB 1;
Pred. No. 1.3e+02;
n: Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 209/035
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                                                                                                                                                                                                                          Length 17;
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US-08-435-628-1114/c
                                RESULT 150
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/373,124
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/986,422
APPLICATION NUMBER: 07/936,422
APPLICATION HOMBER: 07/936,422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Stinchcomb, Dan T.
APPLICANT: Draper, Kenneth
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C.
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   . ZIP: 90071
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Warburg, Richard REGISTRATION NUMBER: 32, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/01 FILING DATE: 05-MAY-1995 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
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STATE: California
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                                                                                                                                                              1584 CATGAACTCCAACACC 1599
                                                                                                                             1 CAAGAACUCCUACACC 16
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633 West Fifth Street
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                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                               Length 17;
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Sequence 1114, Application US/08435628 Patent No. 5817796

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GENERAL INFORMATION:

APPLICANT:

Draper, Kenneth McSwiggen, James Jarvis, Thale

Stinchcomb, Dan T

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US-08-435-628-1116/c
; Sequence 1116, Application US/08435628
; Patent No. 5817796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   밁
                                                                                                                                                                                                                                                                                                                                                                                   RESULT 151
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query match 0.9%;
Best Local Similarity 87.5%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                  GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                              APPLICANT:
APPLICANT:
APPLICANT:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                            TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/0:
FILING DATE: 05-MAY-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM COMPUTER: IBM P.C.
OPERATING SYSTEM: IBM P.C.
OPERATING SYSTEM: IBM P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX:
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FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/245,466
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STATE: California
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                                                                                                                                                                                                                                                                          INFORMATION:
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nucleic acid
                                                                    I: McSwiggen, James
I: Jarvis, Thale
INVENTION: METHODS AND COMPOSITIONS FOR
INVENTION: TREATMENT OF RESTENOSIS AND
INVENTION: CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Suite 4700
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633 West Fifth Street
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                                                                                                                                                                                                                  Draper, Kenneth
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linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (213) 955-0440
                                                                                                                                                                                                                                              Stinchcomb, Dan T.
Lyon & Lyon
                               ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              January 13, 1995
NUMBER: 08/245,466
May 18, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TREATMENT OF RESTENOSIS AND CANCER USING RIBOZYMES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2627
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 12.8; DB 1;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOS 5.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                 Patent No. 5817796
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INFORMATION FOR SEQ ID NO:
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APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
FILING DATE: August 26, 1992
ATTORNEY/AGENT INFORMATION:
          STREET: Sulle
STREET: Los Angeles
CTATE: California
"I S.A."
                                                                                                                                            APPLICANT: McSwiggen, James
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHODS
TITLE OF INVENTION: TREATMEN
TITLE OF INVENTION: CANCER L
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
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APPLICATION NUMBER:
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APPLICATION NUMBER: US
FILING DATE: 05-MAY-19
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                                                                                                                                                                                                                                  APPLICANT: Stinchcomb, Dan APPLICANT: Draper, Kenneth
                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                  NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                               0.9%;
Local Similarity 87.5%;
les 14; Conservat:...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
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REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                               ADDRESSEE:
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90071
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                                                              Suite 4700
                                                                            E: Lyon & Lyon
633 West Fifth Street
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linear
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                                                                                                                                                                                                                                                   Stinchcomb, Dan T.
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                                                                                                                                                METHODS AND COMPOSITIONS FOR TREATMENT OF RESTENOSIS AND CANCER USING RIBOZYMES
                                                                                                                                2627
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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COMPUTER READABLE FORM:

TYPE: 3.5" Diskette, 1.44 Mb

MEDIUM TYPE:

storage

CURRENT APPLICATION DATA:

Word Perfect 5.1 IBM Compatible

IBM P.C. DOS 5.0

OPERATING SYSTEM:

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Matches
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Patent No. 5821046
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                                                                OPERATING SYSTEM: PC-DOS/MS
SOFTWARE: WORDERFECT 6.1
CURRENT APPLICATION DATA:
APPLICATION UMMER: US/08/5
FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
07/960
                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
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FILING DATE: May 18, 1995
APPLICATION NUMBER: 08/249,466
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7 1001
APPLICATION February 7 1001
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathleen M.
                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall CTITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: August 26, 1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER: 08/373,124
FILING DATE: January 13, 1995
                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                            CIT: MA
STATE: MA
TO: 02111
                                                                                                                                                                 MEDIUM TYPE: Diskette, 3.50 inch, COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: February 7
APPLICATION NUMBER: 07/
FILING DATE: December 7
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                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                            CITY: Boston
                                                                                                                                                                                                                                                                                               STREET:
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                                                                                                                                                                                                                                                                                                 One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                         03/19/93
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ber 7, 1992
                                                         07/960,370
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                                                                                                                US/08/541,950B
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US-08-541-950B-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19, Application US/08541950B Patent No. 5821046 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                          TELEFAX: (617) 345-9111
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 bases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 02111
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/541,5
FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/960,370
FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C
TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
NUMBER OF SEQUENCES: 26
                                                                                     TOPOLOGY: linear MOLECULE TYPE: synthetic
                                                                                                                                                                                                                                   TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
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MOLECULE TYPE: synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cli.
STATE: Pu.
CO. 02111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 17 bases
TYPE: nucleic acid
STRANDENNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatib
OPERATING SYSTEM: PC-DOS/
SOFTWARE: WordPerfect 6.1
           LOCATION: 10 -
OTHER INFORMATION: N is 2'-deoxythymidine
                                                                                                                                                                                                                                                                     NAME: Williams, Ph.D., Kath
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
LOCATION: 9
OTHER INFORMATION: N is 2'-deoxythymidine
                                             NAME/KEY: misc_feature
                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Banner & Witcoff, Ltd.
STREET: One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1490 AGCCAGACTTCAGCAGC 1506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 AGCCAGAUNUGAGCAGC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (617) 345-9111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diskette, 3.50 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.9%; Score 12.8; 76.5%; Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                      US/08/541,950B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA
                                                                                                                                                                                                                                                                                                             Kathleen M.
                                                                                                                                                                                                                                                                         3777/57347 (MRC-011AX)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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Score 12.8;

DB 1;

Length 17;

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US-08-541-950B-21
                                                                                                                                                                   US-08-541-950B-22
                                                                                                                                                                                                                                                                                                                                                                                               US-08-541-950B-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPAX: (617)
TELEPAX: (617)
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                         Sequence 22, Application US/08541950B Patent No. 5821046 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Karn J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/960,37
FILING DATE: 03/19/93
ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathle
REGISTRATION NUMBER: 34,380
                                 NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                        APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 3777/57347 (MRC-011AX)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
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ADDRESSEE: Banner & Witcoff, Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: 9 - OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC OPERATING SYSTEM:
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STREET: One Financial Center, 45th Floor
                     DDRESSEE:
                                                                                                                                                                                                                                                                                1490 AGCCAGACTTCAGCAGC 1506
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02111
                                                                                                                                                                                                                                                                                                                                         Similarity
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                 Banner & Witcoff, Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         synthetic RNA
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                                                                                                                                                                                                                                                                                                                                       0.9%;
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                                                                                                                                                                                                                                                                                                                                     Score 12.8; DB 1; Length 17; Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.44 Mb storage
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                                                                                                                                                                                                                                                                                                                     0;
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US-08-628-145-23/c
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              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAECHTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/628,145
FILING DATE: 04-APR-1996
CLASSIFICATION DATA:
APPLICATION UNMBER: US/08/179,738
APPLICATION NUMBER: US/08/179,738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 23, Application US/08628145 Patent No. 5872214
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TELEFAX: (617) 345-9111
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/541,95
FILING DATE: 10/10/95
PRIOR APPLICATION DATA:
APPLICATION UMBER: 07/960,370
FILING DATE: 03/19/93
ATTORNEY/ACENT INFORMATION:
ANAME: MILITED PARTY PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Seizinger, Bernd R. APPLICANT: Kley, Nikolai A. APPLICANT: Bianchi, Albert B.
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LENGTH: 17 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Williams, Ph.D., Kath
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
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02111
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CITY: Palo Alto
STATE: Californ:
                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: UZIP: 94301
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1490 AGCCAGACTTCAGCAGC 1506
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13; Conservative
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SYSTEM: PC-DOS/MS-DOS
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LO-JAN-1994
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76.5%;
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26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.50 inch, 1.44
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Pred. No. 1.3e+02;
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US-08-985-162-184/c
US-08-985-162-184
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Matches 14; Conservative
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INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 18
SEQUENCE CHARACTERISTICS:
                                                                                                                                                   NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 230
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/036,4 FILING DATE: 31 January 1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/985,162
FILING DATE: 04 December 1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Fast-SEO for Windows 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Akhtar, Saghir
APPLICANT: Fell, Patricia
APPLICANT: McSwiggen, James
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 617-8999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ELECTIONS: (415)
            LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
TOPOLOGY: lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1579 TCCTCCATGAACTCCA 1594
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OF DISEASES OR CONDITIONS RELATED
TO LEVELS OF EPIDERWAL GROWTH
FACTOR RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1877
                                                                                                                                                                                                                                                                               60/036,476
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                                                                                                      184:
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Pred. No. 1.3e+02;
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US-08-985-162-326
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                                                                  Best Local Similarity
Matches 9; Conserv
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                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: IBM P.C. DOS 5.0
SOPTWARE: PASTISEO for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/985,162
FILING DATE: 04 December 1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 60/036,476
FILING DATE: 31 January 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                      TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                       NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE DOCKET NUMBER: 23
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                  LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                        TELEPHONE: (213) 955-0440
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STREET: 633 West Fifth Street
STREET: Suite 4700
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STATE: California
                                 2323 AGTGATGTCTGGTCCT 2338
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AGUGAUGUCUGGAGCU 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IBM Compatible
                                                                                                                                                                                                                                                                                        (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.9%;
                                                                  0.9%; Score 12.8; DB 1; 56.2%; Pred. No. 1.3e+02; tive 5; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENZYMATIC NUCLEIC ACID TREATMENT
OF DISEASES OR CONDITIONS RELATED
TO LEVELS OF EPIDERMAL GROWTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FACTOR RECEPTORS
                                                                                                                                                                                                                                                                                                                           230/107
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Pred. No. 1.
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                                                                                                  Length 17;
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RESULT 160 US-09-083-756A-18

Sequence 18, Application US/09083756A Patent No. 6114109

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RESULT 161
US-09-083-756A-19
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                                                                                                                                                                                                                                                                                                                                          Sequence 19, Application US/09083756A Patent No. 6114109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (617) 345-91
TELEFAX: (617) 345-91:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Williams, Ph.D., Kathl
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 377
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                                                                                                                                                                                        APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                     SOFTWARE: WordPerfect 6.1 CURRENT APPLICATION DATA:
                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 incl
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 17 bases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/541,950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
PRIOR APPLICATION DATA
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STATE: ...
To: 02111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Diskette, 3.50 inch, COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: 9
OTHER INFORMATION: N is 2'-deoxythymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature Inocation: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                           INFORMATION:
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                                                                                                                                                                                               Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%;
Similarity 76.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
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                                                                                                                                                                                                                   One Financial Center, 45th Floor
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One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (617) 345-9111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss: single
linear
                                                                                                                                                                                                                                    Banner & Witcoff, Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    synthetic RNA
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                                     US/09/083,756A
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                                                                                                                            3.50 inch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 12.8; DB 1;
Pred. No. 1.3e+02;
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                                                                                                                            1.44 Mb storage
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US-09-083-756A-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21, Application US/09083756A Patent No. 6114109
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (617) 345-9111 INFORMATION FOR SEQ ID NO:
                                                                                                                                                         TELEFAX: (617) 345-91 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                     ATTORNEY AGENT INFORMATION:
NAME: Williams, Ph.D., Kathl
NAME: Williams, 34,380
REGISTRATION NUMBER: 34,380
                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/083,7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 17 bases
                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
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ATTORNEY/AGENT INFORMATION:
                                               MOLECULE TYPE:
                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C
TITLE OF INVENTION: VIRAL (HIV) GROWTH INHIBITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: PL. 02111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Williams, Ph.D., Kath
REGISTRATION NUMBER: 34,380
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LOCATION:
                  NAME/KEY:
                                                                                   STRANDEDNESS:
                                                                                                                                                                                          TELEPHONE:
                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1490 AGCCAGACTTCAGCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 AGCCAGAUUNGAGCAGC 17
                                                                                                     nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Banner & Witcoff, Ltd.
One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17 bases
                                                                                                                                                                        : (617) 345-91
(617) 345-9111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
              misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Diskette,
                                                    synthetic RNA
                                                                                   single
                                                                                                                                                                                          345-9100
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                                                                                                                                                                                                                                                                                                                                                                    US/09/083,756A
                                                                                                                                                                                                                                                                                                                   08/541,950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19:
                                                                                                                                                                                                                                                               Kathleen M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.50 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kathleen M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
                                                                                                                                                                                                                                3777/57347 (MRC-011AX)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3777/57347 (MRC-011AX)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12.8; DB 1;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 17,
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RESULT 164
US-08-584-040-1992/c
"---ance 1992, Application US/08584040
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                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity Matches 13; Conserv
                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22, Application US/09083756A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (617) 345-91:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Williams, Ph.D., Kathle
REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 3777
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 345-9100
TELEFAX: (617) 345-9111
                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff, Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Karn J, Gait MJ, Heaphy S, Dingwall C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Diskette, 3.50 inch, COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                        LOCATION: 10
OTHER INFORMATION: N is 5-bromo-2'-deoxyuridine
                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                     1490 AGCCAGACTTCAGCAGC 1506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1490 AGCCAGACTTCAGCAGC 1506
                                                                                                                                                                                                      13;
                                                                                                                                          1 AGCCAGAUUNGAGCAGC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02111
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                                                                                                                                                                                                                                                                                                                                                                                                     17 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        One Financial Center, 45th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                         synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.9%;
76.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08/541,950
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kathleen M.
                                                                                                                                                                                                   Score 12.8; D
Pred. No. 1.3e
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3777/57347 (MRC-Ollax)
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Pred. No. 1.
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1.3e+02;
3;
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                                                                                                                                                                                                                               DB 1; Length 17;
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ADDRESSEE:

633 West Fifth Street

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RESULT 165
US-08-584-040-2009
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                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                    Patent No.
                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Stinchcomb APPLICANT: Escobedo, TITLE OF INVENTION: 7 TITLE OF SEQUENCES:
                      TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: TREATMENT OF DISEASES OR TITLE OF INVENTION: CONDITIONS RELATED TO LEVEL TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL TITLE OF INVENTION: GROWTH FACTION
                                                                                                                          APPLICANT: Pavco, rames
APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: October 26, 1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (213) 489-160
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 633 West Fifth
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 90071-2066
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                                                                                                                                                                                                                                                                                                                                                                  1909 AATATCATAAATCTTC 1924
                                                                                                                                                                               INFORMATION:
CANT: Pavco, Pamela
                                                                                                                                                                                                                                     2009,
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                                                                                                                                                                                                                      6346398
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Escobedo, Jaime
                                                                                                                                                                                                                                   Application US/08584040
                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                      0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             METHOD AND REAGENT FOR TREATMENT OF DISEASES OF CONDITIONS RELATED TO LI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GROWTH FACTOR
                    8502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   218/064
                                                                                                                                                                                                                                                                                                                                                                                                                      Score 12.8;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                    1.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                           LEVELS
                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: 67-3510
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Pavco, rumanes
APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: October 26, 1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                   UMBER OF SEQUENCES:
                                                                                                                                                                       ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (213) 955-0440
                                               COMPUTER: IBM CONCOPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
les 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A. ZIP: 90071-2066
                                SOFTWARE:
                                                                                                                                        COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/005,974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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California
                                                                                                                                                                                                                                                                                                                  INVENTION: METHOD AND REAGENT FOR THE INVENTION: TREATMENT OF DISEASES OR INVENTION: CONDITIONS RELATED TO LEVELS INVENTION: OF VASCULAR ENDOTHELIAL
                           Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/08584040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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                                                                   IBM Compatible
                DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.9%; Score 12.8; DB 1; Length 17; 75.0%; Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM P.C.
                                                                                                                                                                                                                                                                                                   GROWTH FACTOR
                                                                                                                                                                                                                                                                                   8502
US/08/584,040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        218/064
                                                   DOS 5.0
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US-08-584-040-4140
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                                                                                                                                                                                                                        ZIP: 9071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (213) 955-04
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 211
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: October 26, ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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STRANDEDNESS: &...
TOTY: linear
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                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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Escobedo, Jaime
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87.5%;
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Pred. No. 1.
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Query Match 0.9%;
Best Local Similarity 50.0%;
Matches 8; Conservative
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                                                                                                                                                           TELEFAX: (213) 955-044
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UMBER: 60/005,974
FILING DATE: October 26, 1995
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: McSwiger, James
APPLICANT: McSwiger, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Becobedo, Jaime
TITLE OF INVENTION: MCSWIGHT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR EMDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 8
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                                                                                                                                                                                                     TELEPHONE: (213) 955-0440
                                                                                           STRANDEDNESS: 811
TOPOLOGY: linear
                                                                                                                          LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U.S.A. ZIP: 90071-2066
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Local Similarity 68.8%;
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION:
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': Los Angeles
E: California
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633 West Fifth Street
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January 11, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8502
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pred. No. 1.3e+02;
3; Mismatches 2;
Score 12.8; DB 1;
Pred. No. 1.3e+02;
6; Mismatches · 2;
                                    Length 17;
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Indels
0
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Gaps
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                                                                   RESULT 170
US-08-584-040-8106/c
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                 Sequence 8106, Application US/08584040 Patent No. 6346398 GENERAL INFORMATION:
                                                                                                                                                                                                                        Matches
APPLICANT: Pavco, Pamela
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TELEFAX: (22...
TELEFAX: 67-3510
TELEX: 67-3510
TELEX: 67-3510
TOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                 TOPOLOGY:
US-08-584-040-7971
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COMPUTER: IBM P.C. DOS 5.0
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974

                                                                                                                                                                         Query Match
Best Local Similarity
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Suite 4700
CITY: Los Angeles
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STREET: 63
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                                            1843 ACAGAGAAAGACCTTT 1858
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16 ACAGAGAAACCCCTTT 1
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5. 6346398
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633 West Fifth Street
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                                                                                                                                            Conservative
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Stinchcomb, Dan T.
Escobedo, Jaime
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                                                                                                                                                                         0.9%;
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                                                                                                                                                                         Score 12.8;
Pred. No. 1.
                                                                                                                                            Mismatches
                                                                                                                                                                      1.3e+02;
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                                                                                                                                                                                                              Sequence 21, Application US/09340861
Patent No. 6432704
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 210
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                 TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/00 FILING DATE: October 26, ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM
                                                                                                                CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/584,040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Com
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
STATE: California
                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
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633 West Fifth Street
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                                                                                   611 West Sixth Street
                                                                                                                                                                                                  Kenneth G. Draper
                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 12.8; DB 1; Length 17; larity 87.5%; Pred. No. 1.3e+02; Conservative 0; Mismatches 2; Indels
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Escobedo, Jaime
ENTION: METHOD AND REAGENT FOR THE
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                                                                                                                                                   VIRUS REPLICATION
                                                                                                                                                                METHOD AND REAGENT FOR INHIBITING HERPES SIMPLEX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IBM P.C. DOS 5.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21, Application US/09634262
Patent No. 6440719
GENERAL INFORMATION:
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Best Local (
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 07/882,921
FILING DATE: May 14, 1992
APPLICATION NUMBER: 07/948,359
FILING DATE: September 18, 1992
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
FILING DATE: 07/882,921
APPLICATION NUMBER: 07/882,921
FILING DATE: May 14, 1992
APPLICATION NUMBER: 07/948,359
FILING DATE: September 18, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                    STREET: 611 West :
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: IBM P.C. DOS (Version 5.0) SOFTWARE: WordPerfect (Version 5.1)
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mes 12; Conserv
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TOPOLOGY: 111
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                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE:
                                                                                                      APPLICATION NUMBER:
                                                                                                                                                      FILING DATE:
                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
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                                                                                                                                                                                                                                                                                                            USA
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Pred. No. 1.3e+02;
2; Mismatches 2
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US-09-474-432B-783
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APPLICANT: EL SOLH, NEVINE
APPLICANT: EL SOLH, NEVINE
APPLICANT: ALLIGNET, JEANINE
APPLICANT: ALLIGNET, JEANINE
TITLE OF INVENTION: POLYNUCLEOTIDES AND THEIR USE FOR DETECTING RESISTANCE
TITLE OF INVENTION: COMPOUNDS
TILE REFERENCE: 03715-0059
CURRENT APPLICATION UNMEER: US/09/446,301A
CURRENT FILING DATE: 1999-12-20
NUMBER OF SEQ ID NOS: 51
SEQ ID NO 31
LENGTH: 17
LENGTH: 17
LENGTH: 17
LENGTH: 17
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                                                                                                                                                                                                                        Sequence 783, Application US/09474432B Patent No. 6528640
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                                                                                        APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                             APPLICANT: Ribozyme Pharmaceuticals, APPLICANT: Beigelman, Leo APPLICANT: Burgin, Alex
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TELEPAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: :
               APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleotide triphosphate and their incorporation into oligonucleot
FILE REFERENCE: MBHB00-831-B (247/276)
                                                      APPLICANT:
CURRENT APPLICATION NUMBER: US/09/474,432B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 201
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
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                                                                   Beaudry, Amber
Karpeisky, Alex
Adamic, Jasenka
Sweedler, David
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Pred. No. 1.3e+02;
2; Mismatches 2
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; Sequence 554, Application US/09371772B
; Patent No. 6566127
; GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
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US-09-371-772B-537/c
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; ORGANISM: Homo sapiens
US-09-474-432B-783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR FILING DATE: 1997-11-05
PRIOR APPLICATION NUMBER: US 60/084,727
PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR FILING DATE: 1998-11-04
PRIOR FILING DATE: 1998-11-04
PRIOR APPLICATION NUMBER: US 09/301,511
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Best Local S
Matches 14
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SEQ ID NO 537
LENGTH: 17
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GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
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Best Local Similarity
Matches 10; Conservat
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 80/584,040
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Homo sapiens
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les 14; Conserv
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McSwiggen, Jim
Stinchcomb, Dan
Escobedo, Jaime
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87.5%; Pred. No. 1.
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Pred. No. 1.
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1.3e+02;
thes 2; Indels
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1.3e+02;
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TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

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RESULT 178
US-09-371-772B-1907
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; ORGANISM: Homo sapiens
US-09-371-772B-554
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                                               GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals,
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
                                                                                                                                                                                    Sequence 1907, Application US/09371772B Patent No. 6566127
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Best Local (
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Patent No. 6566127
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Best Local
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PRIOR PILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
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APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
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CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Homo sapiens
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; ORGANISM: Mus sp.
US-09-371-772B-2681
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
                                                                                                                                                                        GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PILING DATE: 1995-10-26
PRIOR PILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
                                                                              APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: "Stinchcomb, Dan
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CURRENT APPLICATION UNMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR PPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION SUMBER: US 08/584,040
PRIOR PILING DATE: 1996-01-08
PRIOR PILING DATE: 1996-01-08
                 APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
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FILE REFERENCE: MBHB00,876-J
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Local Similarity 68.8%;
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Pred. No. 1
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Pred. No. 1.3e+02;
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RESULT 182
US-09-371-772B-4565/c
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; ORGANISM: Mus sp.
US-09-371-772B-3889
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                                                                                                                                                                                                                                    Sequence 4565, Application US/09371772B Patent No. 6566127
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Best Local Similarity
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LENGTH: 17
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APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions |
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TILE REFERENCE: MBHB00,876-J (237/198)
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
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APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Scobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of (237/198)
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PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                        APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
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SOFTWARE: PatentIn version 3.0
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Pred. No. 1.
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Pred. No. 1.3e+02
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RESULT 184
US-09-371-772B-6727
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US-09-371-772B-5571/c
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                                                                                                                         GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam

APPLICANT: McSwiggen, Jim

APPLICANT: Stinchcomb, Dan

APPLICANT: Escobedo, Jaime.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 5571
LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                       Sequence 6727, Application US/09371772B Patent No. 6566127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.9%;
Best Local Similarity 87.5%;
Matches 14; Conservative
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APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MEMBBO0,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
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APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions:
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
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APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
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PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
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Local Similarity 87.5%;
les 14; Conservation
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Pred. No. 1.
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Pred. No. 1.
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; ORGANISM: Homo sapiens
US-09-371-772B-6727
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                                                                                                                                                                                                                                                                         Sequence 6762, Application US/09371772B Patent No. 6566127
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                APPLICANT: Pavcó, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MeBHB00, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MedhBo0, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
CURRENT FILING DATE: 1999-08-10
                                                                                                                                  APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
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SOFTWARE: PatentIn version 3.0
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PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
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PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
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APPLICANT: Pavco, Pam
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TYPE: RNA
ORGANISM: Homo sapiens
PRIOR APPLICATION NUMBER: US 60/005,974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH: 17
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11 Similarity 81.2%;
13; Conservation
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Similarity 56.2%;
9; Conservation
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Pred. No. 1.3e+(
5; Mismatches
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Pred. No. 1.3e+02;
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; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-371-772B-6762
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US-09-099-932-29/c
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                                                                                                                                                                                                                                                                    Sequence 29, Application US/09099932 Patent No. 6570001 GENERAL INFORMATION:
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Best Local Similarity
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         APPLICANT: E1 Solh, Nevine
APPLICANT: Allignet, Jeanine
APPLICANT: Allignet, Jeanine
ITILE OF INVENTION: POLYMUCLEOTIDES AND THEIR USE FOR DETECTING RESISTANCE
ITILE OF INVENTION: CONCOUNDS
ITILE OF INVENTION: CONCOUNDS
FILE REFERENCE: 03495.0173-0000
CURRENT APPLICATION NUMBER: US/09/099,932
CURRENT APPLICATION NUMBER: US/09/099,932
CURRENT FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: 60/050,380
EARLIER FILING DATE: 1997-06-20
NUMBER OF SEQ ID NOS: 50
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APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE REFERENCE: MBHB00,876-J (237/198)
CURRENT REPERENCE: MBHB00,876-J (237/198)
CURRENT FILLING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
RUGHER PRIOR SEQ ID NOS: 14225
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APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
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PRIOR APPLICATION NUMBER: US
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn Ver.
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ORGANISM: Homo sapiens
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12.8; DB 1;
Pred. No. 1.3e+02;
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Pred. No. 1
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US-08-651-472-75
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                                                                                                                                                                                                                                       APPLICATION NUMBER: US 07/750,080
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 30472/166/
TELEPHONE: (202)672-5300
TELEPHONE: (202)672-5399
                                                                                                                                                                                                          NFORMATION FOR SEQ ID NO:
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APPLICANT:
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APPLICANT: PFLEIDERER, Michael
TITLE OF INVENTION: DIRECT MOLECU
TITLE OF INVENTION: VIRUSES CONTA
TITLE OF INVENTION: (HIV-1) ANTIG
                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 14 base pairs
                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US 07/914,738
FILING DATE: 20-JUL-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                 TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid;
DESCRIPTION: Synthetic DNA oligonucleotide
IMMEDIATE SOURCE:
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NAME/KEY:
LOCATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
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                                                                                                                                     STRANDEDNESS:
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                                                                                                                                                  nucleic acid
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                                                  gp160 in vselP-gp160 virus
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CDS
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Pred. No. 1.3e+02
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RESULT 191
US-08-358-928-75
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CURRENT FILING DATE: 1997-12-24
EARLIER APPLICATION NUMBER: 60/037,658
EARLIER FILING DATE: 1997-01-23
EARLIER APPLICATION NUMBER: 08/373,124
EARLIER FILING DATE: 1995-01-13
EARLIER APPLICATION NUMBER: 08/245,466
EARLIER FILING DATE: 1994-05-18
NUMBER OF SEQ ID NOS: 375
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: JARVIS, THALE
APPLICANT: MCSWIGGEN, JAMES A.
APPLICANT: STINCHCOMB, DAN T.
APPLICANT: STINCHCOMB, DAN T.
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES
TITLE OF INVENTION: OR CONDITIONS RELATED TO LEVELS OF C-FOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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COUNTRY: USA
ZIP: 20007-5109
ZIP: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No. 6103890
                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                      APPLICANT: FALKNER, FAIKO GUNTEET
APPLICANT: PFLEIDERER, Michael
TITLE OF INVENTION: DIRECT MOLECULAR CLONING OF CHIMERIC
TITLE OF INVENTION: VIRUSES CONTAINING HUMAN IMMUNODEFCIENCY VIRUS TYPE
TITLE OF INVENTION: (HIV-1) ANTIGENS
NUMBER OF SEQUENCES: 95
                                                                                                                                                                                                                                                                                                                           APPLICANT: SCHEIFLINGER, Friedrich APPLICANT: FALKNER, Falko Gunter APPLICANT: PFLEIDERER, Michael
                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                            CITY: Washington
                                                                                                                                                                               ADDRESSEE: Foley & Lardner STREET: 3000 K Street, N.W
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Pred. No. 1e+02;
0; Mismatches 1;
                                                                                                                                                                                   Suite 500
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      Version #1.30
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Best Local
                                                                 Matches
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                                                                                            Query Match
                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/535,366C CURRENT FILING DATE: 2000-03-24 PRIOR APPLICATION NUMBER: 60/125,864 PRIOR FILING DATE: 1999-03-24
                                                                                                                                                                                                                                                                                                                                                            APPLICANT: SYKES, KATHRYN F.
APPLICANT: JOHNSTON, STEDHEN A.
TITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS
FILE REFERENCE: UTSD:557
                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 5
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                             LENGTH: 14
TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                               OTHER INFORMATION: OTHER INFORMATION:
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FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (202)672-5300
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DESCRIPTION: Synthetic DNA oligonucleotide
                                                                                                                                                                                                                                                ID NO 3
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                                                               Similarity 9; Conserv
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                                                               Conservative
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                                                                                                                                                Description of Artificial Sequence: Primer
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                                                              Score 12.4; DB 1; Length 14; Pred. No. 1e+02; 4; Mismatches 1; Indels
                                                                                                                                                               Synthetic
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                                                               Gaps
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US-09-535-366C-5/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 5, Application US/09535366C Patent No. 6410241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: SYKES, KATHRYN F.
APPLICANT: JOHNSTON, STEPHEN A.
TITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS
FILE REFERENCE: UTSD:557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/535,366C
CURRENT FILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: 60/125,864
PRIOR FILING DATE: 1999-03-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/334,847
FILING DATE: No. 5693532ember 4, 1994
              ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,:
REFERENCE/DOCKET NUMBER:
                                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: METHOD AND REAGTITLE OF INVENTION: INHIBITING RESPONDED OF INVENTION: SYNCYTIAL VIRUS
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                      STREET: 633 West Fi
STREET: Suite 4700
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
nes 13; Conserv
                                                                                                                                                                                                           COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                       California
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633 West Fifth Street
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                                                                                                                                                                                              Word Perfect 5.1
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                             32,327
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Pred. No. 16
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US-08-363-240A-63/c ; Sequence 63, Application US/08363240A ; Patent No. 5705388
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                                                                               US-08-363-240A-63
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base ~~'
Type
0.9%;
Best Local Similarity 92.9%;
Matches 13; Conservative
                                                                                              TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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Best Local Similarity 71.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                     TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/363,
FILING DATE: December 23, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: METHOD AND REAGENT FOR TITLE OF INVENTION: PREVENTION, INHIBITION TITLE OF INVENTION: PROGRESSION AND RECRESTITLE OF INVENTION: OF VASCULAR DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
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CITY: Los Angeles
STATE: California
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                      NAME: Warburg, Richard REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
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Bisgaier, Charles
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          Score 12.4; DB 1; Length 15; Pred. No. 1.2e+02; 0; Mismatches 1; Indels
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Pred. No. 1.2e+02;
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US-08-657-884-27/c
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Patent No. 5858981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local :
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                                                                                                                                                                                                                                           GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
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                                                                     CORRESPONDENCE ADDRESS:
ADDRESSE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
CITY: ARLINGTON
STATE: VIRGINIA
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                APPLICANT: SCHREIBER, ALAN D.
APPLICANT: PARK, JONG-GU
TITLE OF INVENTION: METHODS OF INHIBITING PHAGOCYTOSIS
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/657,884 FILING DATE: 07-JUN-1996
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 55
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 816-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
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TITLE OF INVENTION: METHODS O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA (genomic)
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                                      COUNTRY: U.S.A. ZIP: 22201-4714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TORNEY/AGENT J.
NAME: WILSON, MARY J.
32,955
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                             1783 GACAAAGACAAGCC 1796
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1100 NORTH GLEBE ROAD, 8TH FLOOR
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---- TD NO: 27:
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Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 12.4; DB 1; 92.9%; Pred. No. 1.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-585-684B-2115/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; MOLECULE TYPE: DNA (genomic) US-08-657-884-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (703) 816-4100 INFORMATION FOR SEQ ID NO: 31:
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INFORMATION FOR SEQ ID NO: 2115:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                              NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                        APPLICATION NUMBER: 60/000,951 FILING DATE: July 7, 1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                             APPLICATION NUMBER: US/08/585,684B FILING DATE: January 16, 1996 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Jarvis, Thale APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JLECOMMULE: (/UL)
TELEPHONE: (/UL)
TELEPHONE: (703) /
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REGISTRATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/657,884
FILING DATE: 07-JUN-1996
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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o. 5877021
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                                                                                                                                                                                                                                                                                                                                                                                  90071
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         633 West Fifth Street
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                                                                                                                                                                                                                                                                                                                                                                                                 U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             METHOD AND REAGENT FOR THE INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12.4; DB 1; Length 15; Pred. No. 1.2e+02; 0; Mismatches 1; Indels
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US-08-716-308-13
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                                                           Query Match
Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/716,308
FILING DATE: 24-SEP-1996
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9405951.6
FILING DATE: 25-MAR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/C
FILING DATE: 27-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                    NAME: Hohenschutz, Liza D.
REGISTRATION NUMBER: 33,712
REFERENCE/DOCKET NUMBER: PPD40027X/UST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 1800 Conc
CITY: Wilmington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Match 0.9%;
Local Similarity 92.9%;
les 13; Conservative
                                                                                                                                                                                      TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 19850
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                            2166 TGTTTTGGTAACAG 2179
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 2 TGTTTTCGTAACAG 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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                                                              Conservative
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                                                                                                                                                                                                                                                      (302) 886-1699
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                                                                             Score 12.4; DB 1;
Pred. No. 1.2e+02;
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Pred. No. 1.2e+02;
0; Mismatches 1;
                                                              Mismatches
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                                                                                              Length 15;
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RESULT 200 US-08-913-833-157/c ; Sequence 157, Application US/08913833

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RESULT 201
US-09-038-073-2115/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: EP 96870005.4
FILING DATE: 26 Jan 1996
PRIOR APPLICATION DATA:
APPLICATION UNMBER: EP 96870081.5
FILING DATE: 25 Jun 1996
ATTORNEY/AGENT INFORMATION:
NAME: KAMMERER, PATRICIA A.
REGISTRATION NUMBER: 29,775
REFERENCE/DOCKET NUMBER: INNS:008
INFORMATION FOR SEQ ID NO: 157:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                         Sequence 2115, Application US/09038073 Patent No. 6194150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.9%; Score 12.4; Best Local Similarity 92.9%; Pred. No. 1.
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APPLICATION NUMBER: US/08/913,833 FILING DATE: 15 Sep 1997 FILING APPLICATION DATA:
APPLICATION NUMBER: PCT/EP97/00211 FILING DATE: 17 Jan 1997 PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                       GENERAL INFORMATION:
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ZIP: 77210-4433
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                  TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                 APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: DI
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                          CORRESPONDENCE ADDRESS:
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APPLICANT: LOUWAGIE, JOOST
APPLICANT: ROSSAU, RUDI
TITLE OF INVENTION: METHOD FOR DETECTION OF DRUG-INDUCED
TITLE OF INVENTION: MUTATIONS IN THE REVERSE TRANSCRIPTASE GENE
NUMBER OF SEQUENCES: 164
                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: P.O. BOX 4433
CITY: HOUSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                          ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                  1856 TTTCTGATCTGGTG 1869
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               E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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US-09-158-980-27/c
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Patent No. 6242427
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/158,980
FILING DATE:
                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: IBM P.C. DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
ATTORNEY/AGENT INFORMATION:
NAME: WILSON, MARY J
REGISTRATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 55
                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: SCHREIBER, ALA APPLICANT: PARK, JONG-GU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/038,073 FILING DATE: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: METHODS OF INHIBITING PHAGOCYTOSIS NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A. ZIP: 22201-4714
                                                                                                                                                                                                                                                                                                                                               STATE:
                                                                                                                                                                                                                                                                                                                                                             STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: NIXON & VANDERHYE P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                              08/657,884
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   555-46
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Pred. No. 1.2e+02;
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US-09-158-980-31/c
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; MOLECULE TYPE:
US-09-158-980-31
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Best Local Similarity 92.9%;
                                                                      Query Match
Best Local Similarity
                                                          Matches
                                                                                                                                                                                                                                                             FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/657,884
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: WILSON, MARY J.
REGISTRATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 555-46
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DDS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/158,980
                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SCHREIBER, ALAN D. APPLICANT: PARK, JONG-GU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                     TELEPHONE: (703)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
TOPOLOGY: lir
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                           1783 GACAAAGACAAGCC 1796
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                                                                                                                                                                         nucleic acid
GACAAAGACAAGAC 2
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                                                        Conservative
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                                                                                                                                                linear
                                                                                                                                                                                                                                  (703) 816-4000
703) 816-4100
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                                                                                                                               DNA (genomic)
                                                                                                                                                            single
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ID NO: 27:
                                                                      0.9%;
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                                                                                                                                                                                                                         31:
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Pred. No. 1.
                                                                      Score 12.4;
Pred. No. 1.
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                                                                                    Length 15;
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Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/580,794C
CURRENT FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: 08/913,833 now US/6,087,093
PRIOR FILING DATE: 1997-09-15
PRIOR APPLICATION NUMBER: PCT/EP 97/00211
PRIOR FILING DATE: 1997-01-17
PRIOR FILING DATE: 1997-01-17
PRIOR APPLICATION NUMBER: EP 96870081.5
PRIOR APPLICATION NUMBER: EP 96870081.5
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APPLICANT: Louwagie, Joost
APPLICANT: Rossau, Rudi
TITLE OF INVENTION: METHOD FOR DETECTION OF DRUG-INDUCED MUTATIONS IN THE REVERSE
TITLE OF INVENTION: TRANSCRIPTASE GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 1996-06-25
NUMBER OF SEQ ID NOS: 164
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                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial sequence
                                                      OPERATING SYSTEM: IBM P.C. DOS SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                        ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                              APPLICATION NUMBER: FILING DATE: Janua:
                                                                                                     COMPUTER: IBM CON OPERATING SYSTEM:
                                                                                                                                                                                                    COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1856 TTTCTGATCTGGTG 1869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION:
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                                                                                                                       IBM Compatible
....пвык: US/08/584,040
January 11, 1996
DN: 514
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                                                                                                                                                                                                                                                                                                                                                                                                  METHOD AND REAGENT FOR THE TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                                                                                  GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                8502
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Pred. No. 1.2e+02;
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                                                                                                       5.0
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; SEQ ID NO 4128
; LENGTH: 15
; TYPE: RNA
; ORGANISM: Mus sp.
US-09-371-772B-4128
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US-08-527-060-14/c
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US-09-371-772B-4128
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
TELEFAX: (213) 955-0440
TELEFAX: 67-3510
INFORMATION FOR SEQ ID NO: 8473:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                           Sequence 14, Application US/08527060
Patent No. 5934440
GENERAL INFORMATION:
APPLICANT: Goldenberg, Tavi
APPLICANT: Tritz, Richard
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Patent No. 6566127
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MHSH800, 976-J (237/198)
CURRENT FILING DATE: 1995-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PRIOR SEQ ID NOS: 14225
SOFTMARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
     TITLE OF INVENTION:
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%; Score 12.4; DB 1; Local Similarity 71.4%; Pred. No. 1.2e+02; Pres 10; Conservative 3; Mismatches 1
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                                                                                                                                                                                                                                                                                                                   1822 AAGATGTTGAAAGA 1835
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                                                                                                                                                                                                                                                                                                                                                                   10; Conservative
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RIBOZYME THERAPY FOR THE TREATMENT AND/OR PREVENTION OF RESTENOSIS
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APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: McSwiggen, Jim

APPLICANT: Stinchcomb, Dan

APPLICANT: Escobedo, Jaime

FITTE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

FITTE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

FILE REPERENCE: MBHB00,876-J (237/198)

CURRENT APPLICATION NUMBER: US/09/371,772B

CURRENT APPLICATION NUMBER: US/09/371,772B

CURRENT FILING DATE: 1995-08-10

PRIOR APPLICATION NUMBER: US 60/005,974

PRIOR APPLICATION NUMBER: US 08/584,040

PRIOR FILING DATE: 1995-10-26

PRIOR FILING DATE: 1996-01-08

NUMBER OF SEQ ID NOS: 14225

SOFTWARE: PatentIn version 3.0

LENGTH: 16

TYDE: BNA
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                                                                                                                                   ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-371-772B-5827
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US-09-371-772B-5827
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                                            Query Match 0.9
Best Local Similarity 78.6
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 5827, Application US/093717728 Patent No. 6566127
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COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/527,060
FILING DATE: 12-SEP-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: McMasters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 480124.402C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622.4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (206) 682-6031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 6300 C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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1924 CTTGGAGCCTGCAC 1937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14 TTTGGTGACAGAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEED and BERRY
                                       0.9%; Score 12.4; DB 1; 78.6%; Pred. No. 1.3e+02; ative 2; Mismatches 1;
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                                                                                    Length 16;
                                            0;
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                                            Gaps
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US-08-765-340-141

; Sequence 141, Application US/08765340

; Patent No. 6150092
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; Sequence 85, Application US/09475947A
; Patent No. 6472154
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                              ; TYPE: RNA
; ORGANISM: Homo sapiens
US-08-998-099-328
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                                                                                  RESULT 211
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/475,947A
CURRENT FILING DATE: 1999-12-31
NUMBER OF SEQ ID NOS: 346.
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 85
LENGTH: 12
TYPE: DNA
ORGANISM: human
ORGANISM: human
                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 328
LENGTH: 14
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Matches
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                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/08/998,099A
CURRENT FILING DATE: 1997-12-24
EARLIER APPLICATION NUMBER: 60/037,658
EARLIER FILING DATE: 1997-01-23
EARLIER APPLICATION NUMBER: 08/373,124
EARLIER FILING DATE: 1995-01-13
EARLIER APPLICATION NUMBER: 08/245,466
EARLIER APPLICATION NUMBER: 08/245,466
EARLIER APPLICATION NUMBER: 08/245,466
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: JARVIS, THALE
APPLICANT: MCSWIGGEN, JAMES A.
APPLICANT: STINCHCOMB, DAN T.
APPLICANT: STINCHCOMB, DAN T.
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES
TITLE OF INVENTION: OR CONDITIONS RELATED TO LEVELS OF C-FOS
FILE REFERENCE: 231/175
FILE REFERENCE: 231/175
CURRENT APPLICATION NUMBER: US/08/998,099A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No. 6103890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Garner, Harold R.
APPLICANT: Wren, Jonathan D.
APPLICANT: Minna, John D.
TITLE OF INVENTION: Polymorphic Repeats in Human Genes
FILE REFERENCE: UTSD0667
                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 375
                                                                                                                                                                                                          Local 12;
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nes 12; Conserv
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                                                                                                                                                                                                                                    0.9%; Score 12;
100.0%; Pred. No.
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                                                                                                                                                                                                                  1.2e+02;
hes 0;
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                                                                                                                                                                                                                           Sequence 121, Application US/07664989B Patent No. 5223409
                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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COMPUTEN: IBM PC Compatible
COMPUTEN: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V
SOFTWARE: #1.30 (EPO)
CURRENT APPLICATION DATA:
APPLICATION UNMERR: US/08/765,340
FILING DATE: 23-DEC-1996
                                                                                                                                                                                        GENERAL INFORMATION: APPLICANT: Ladner,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (212) 758-4
TELEFAX: (212) 751-684
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 145146/94
EILING DATE: 27-JUN-1994
PRIOR APPLICATION DATA:
APPLICATION UNMBER: JP 311130/94
FILING DATE: 21-NOV-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                               APPLICANT: Markland, William
APPLICANT: Ley, Arthur Charles
APPLICANT: Kent, Rachel Baribault
TITLE OF INVENTION: Directed Evolution of No. 5223409el
TITLE OF INVENTION: Binding Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "synthetic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: SERUNIAN, LESLIE
REGISTRATION NUMBER: 35,353
REFERENCE/DOCKET NUMBER: 14:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: AN ANTISENSE NUCLEIC ACID TITLE OF INVENTION: COMPOUND NUMBER OF SEQUENCES: 185 CORRESPONDENCE ADDRESS:
                   NUMBER OF SEQUENCES: 1
                                                                                                                                                    APPLICANT:
                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 14 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 345 PARK AVENUE CITY: NEW YORK STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 100.0%;
les 12; Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                1935 CACACAGGATGG 1946
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UCHIDA, T.,
TANAKA, Y.,
MATSUDA, Y.,
                                                                                                                                                Guterman, Sonia Kosow
Roberts, Bruce Lindsay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
                                                                                                                                                                                          Ladner, Robert Charles
Browdy and Neimark
                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12;
; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1452-4005
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COUNTRY:

USA

Washington

Suite 300

419 Seventh Street, N.W.

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RESULT 213
US-09-177-359-34/c
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                                                                                                                                                                                                              APPLICANT: SINNETT, Daniel
APPLICANT: LABUDA, Damian
TITLE OF INVENTION: DETECTION OF CYP1A1, CYP2A4, CYP2D6 AND
TITLE OF INVENTION: NATZ VARIANTS BY PCR-ALLELB-SPECIFIC OLIGONUCLEOTIDE (ASO)
TITLE OF INVENTION: ASSAY
FILE REFERENCE: 12667-17"US" FC/1d
CURRENT APPLICATION NUMBER: US/09/177,359B
CURRENT APPLICATION NUMBER: US/09/177,359B
CURRENT FILING DATE: 1998-10-23
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 34
LENGTH: 15
TYPE: DNA
Abidical Communication of Carbon and 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 34, Application US/09177359B Patent No. 6183963
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 100. Matches 12; Conservative
       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/07/664,989B
FILING DATE: 19910301
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US89/03731
PILING DATE: 01-SEP-1989
PRIOR APPLICATION DATA:
APPLICATION UMBER: 07/487,063
PILING DATE: 02-MAR-1990
PRIOR APPLICATION DATA:
APPLICATION UMBER: 07/240,160
PRIOR APPLICATION DATA:
APPLICATION UMBER: 07/240,160
PILING DATE: 02-SEP-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 202-737-3528
INFORMATION FOR SEQ ID NO:
                                                                                                                OTHER INFORMATION:
                                                                                                                                                      FEATURE:
                                                                                                                                                                                ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: doub
TOPOLOGY: circular
MOLECULE TYPE: genom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Cooper, Iver P. REGISTRATION NUMBER: 280 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS_MS-DOS
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                                                                                                            cDNA for use as probes
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       Score 12;
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       DB
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   1;
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Length 15;
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; LENGTH: 15
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-081-646-708
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APPLICANT: Zhou, Wei
APPLICANT: Zhou, Wei
TITLE OF INVENTION: Gene Expression Profiles in
TITLE OF INVENTION: Cancer Cells
FILE REFERENCE: 01107.74646;
CURRENT APPLICATION NUMBER: US/09/081,646
CURRENT FILING DATE: 1998-05-20
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ ID NOS: 871
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-081-646-708
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                                       Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 708
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 708, Application US/09081646 Patent No. 6333152 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 133, Application US/09081646
Patent No. 6333152
GENERAL INFORMATION:
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Best Local :
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Matches 12; Conservative C
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                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Kinzler, Kenneth
APPLICANT: Vogelstein, Bert
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2324 GTGATGTCTGGT 2335
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                                                           Score 12;
Pred. No.
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0; Mismatches 0;
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. 1.3e+02;
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CURRENT APPLICATION NUMBER: US/09/475,947A
CURRENT FILING DATE: 1999-12-31
NUMBER OF SEQ ID NOS: 346
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 180
LENGTH: 15
TYPE: DNA
ORGANISM: human
US-09-475-947A-180
                                                                                                                             RESULT 218
US-08-248-357C-11
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Best Local Similarity
Marches 12; Conserva
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US-09-081-646-835
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; LENOTH: 15
; TYPE: DNA
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-081-646-835
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Best Local S
Matches 12
                                                                                            Patent No.
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                                                                                                          Sequence 11,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Garner, Harold R.
APPLICANT: Wren, Jonathan D.
APPLICANT: Minna, John D.
TITLE OF INVENTION: Polymorphic Repeats in Human Genes
FILE REFERENCE: UTSD0667
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CURRENT FILING DATE: 1998-05-20
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ ID NOS: 871
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                           GENERAL INFORMATION:
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APPLICANT: Kinzler, Kenneth
APPLICANT: Vogelstein, Bert
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APPLICANT: Zhou, Wei
APPLICANT: Luban, Jo
APPLICANT: Goff, Sto
TITLE OF INVENTION:
TITLE OF INVENTION:
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les 12; Conserv
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                                                                                              1, Application US/08248357C 
5773225
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                     Luban, octobre P. Goff, Stephen P.
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1.3e+02;
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; FRAGMENT TYPE: N-terminal
US-08-248-357C-11
CURRENT APPLICATION UNDER: CLASSIFICATION:

2 CLASSIFICATION: 435

ATTORREY/AGENT TO COMPATION: CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                              Sequence 20, Application US/08282197C Patent No. 5871730 GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/248
FILING DATE: 24-MAY-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
                                                                                                                                                                                                                                                                       APPLICANT: Brzezinski, Ryszard
APPLICANT: Dery, Claude V
APPLICANT: Beaulieu, Carole
TITLE OF INVENTION: Thermostable Xylanase DNA, Protein and
TITLE OF INVENTION: Methods of Use
NUMBER OF SEQUENCES: 67
                                                                                                                             ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MEDOS
OPERATING SYSTEM: PC-DOS/MEDOS
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                          STATE: New York COUNTRY: U.S.A. ZIP: 10036
                                                                                                                                                                                           CITY: Washington STATE: DC
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les 12; Conserv
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1185 Avenue of the Americas
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ork Ave., NW
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thes 0;
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                                 Best Local Similarity Matches 12; Conserv
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                                                                   Query Match
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Patent No. 5955266
GENERAL INFORMATION:
APPLICANT: Bray, Paul F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                         TELEFAX: 619/678-5099
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 619/678-5070
                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Haile Ph.D., Liga A.,
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 07:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 4225 Executive Square, Suite 1400
CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Goldschmidt-Clermont, Pascal:
APPLICANT: J.
APPLICANT: J.
TITLE OF INVENTION: USE OF PLATELET POLYMORPHISM PLA2 TO
TITLE OF INVENTION: DIAGNOSE RISK OF THROMBOTIC DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
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TELEPAX: 202-371-2540
                                                                                                                                                       FEATURE:
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                                                                                                                     NAME/KEY:
LOCATION:
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                                                                                                                                                                                                                  TYPE: nucleic acid
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2003 GAGCCCGGAGGC 2014
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                                                                                                                                                                       DNA
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US-09-266-409-8/c
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US-08-626-023-4
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                                                         Patent No.
                                                                                                                                                                                  Sequence 8, Application US/09266409 Patent No. 6225061
                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.9%;
Best Local Similarity 100.0%;
Matches 12; Conservative (
                                                                                                                                                                   GENERAL INFORMATION:
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                                                                     APPLICANT: Becker, Thomas
APPLICANT: Sequenom, Inc.
TITLE OF INVENTION: Systems and Methods for Performing Reactions in an
TITLE OF INVENTION: Environment
FILE REFERENCE: Sequence listing for 24736-2023
CURRENT APPLICATION NUMBER: US/09/266,409
CURRENT FILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Bray, Paul F.
APPLICANT: Goldschmidt-Clermont, Pascal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Haile Ph.D., Lisa A.,
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 0726
TELECOMMUNICATION INFORMATION:
TELECHONE: 619/678-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 619/678-5099 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/626,023 FILING DATE: 01-APR-1996 CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                     2003 GAGCCCGGAGGC 2014
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        619/678-5099
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                                                                                                                                                                                                                                                                                                                                                                                            0.9%; Score 12; DB 1; L
.00.0%; Pred. No. 1.5e+02;
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US-09-678-620-8/c
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                                                                                                                                                                                                                     RESULT 224
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                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Becker, Thomas
APPLICANT: Hubert K"ster
APPLICANT: Charles Cantor
TITLE OF INVENTION: Systems and Methods for Performing Reactions in an Unsealed
TITLE OF INVENTION: Environment
FILE REFERENCE: Sequence listing for performing reactions in an Unsealed
                                                                                                                                                               Sequence 35, Application US/08182968A Patent No. 5610054
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 09/266,409
PRIOR FILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 8
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LENGTH: 16
TYPE: DNA
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                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE:
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LOCATION: (1)..(16)
OTHER INFORMATION: Sequencing primer for exon 7 of human p53
                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: Synth NAME/KEY: primer bind LOCATION: (1)...(16)
OTHER INFORMATION: Sequencing primer for exon 7 of human p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
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                                                                      APPLICANT: Draper, Kenneth G.
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: INHIBITING HEPATITIS C
TITLE OF INVENTION: VIRUS REPLICATION
                                   CORRESPONDENCE ADDRESS:
                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 100.0%; es 12; Conservative (
                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                           1940 AGGATGGGCCTC 1951
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                                                                                                                                                                                                                                                                         AGGATGGGCCTC 1
3: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                              Conservative
                                                         VIRUS REPLICATION
497
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic primer
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; Sequence 301, Application US/08182968A
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Best Local Similarity
Matches 10; Conserv
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GENERAL INFORMATION:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/182
FILING DATE: 13-CANUARY-1994
FRIOR APPLICATION NUMBER: 07/882,88
APPLICATION NUMBER: 07/882,88
EILING DATE: 14-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: WAYDUTG, RICHARD:
REGISTRATION UNMBER: 32,327
REFERENCE/DOCKET NUMBER: 205/
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                 OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/182,968A
                                                                                                                                                      ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3 5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Draper, Kenneth G.
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: INHIBITING HEPATITIS C
TITLE OF INVENTION: VRUS REPLICATION
VRUS REPLICATION
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COMPUTER READABLE FORM:
                              FILING DATE: 13-JANUARY-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                        ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: sing:
TOPOLOGY: linear
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OPERATING SYSTEM:
APPLICATION NUMBER: FILING DATE: 14-MA
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                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                          CITY: Los Angeles
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California
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              07/882,888
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Pred. No. 1.
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RESULT 226
US-08-182-968A-302/c
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Matches
Query Match 0.8%;
Best Local Similarity 86.7%;
Matches 13; Conservative
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                                                                                                                                                                                                       APPLICATION UNMBER: US/08/182,968A
FILLING DATE: 13-JANUARY-1994
PRIOR APPLICATION DATA: 07/882,888
FILLING DATE: 14-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 205/277
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
                                                                                                                                                                          TELEFAX: (213) 955-04
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Draper
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INFORMATION FOR SEQ ID NO:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Draper, Kenneth G.
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: INHIBITING HEPATITIS C
TITLE OF INVENTION: VIRUS REPLICATION
NUMBER OF SEQUENCES: 497
                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                   TYPE: nucleic acid
STRANDEDNESS: singl
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 86.7
nes 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (213) 489-16
TELEFAX: (213) 955-0440
                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3: Lyon & Lyon
633 West Fifth Street
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                                                                                                       single
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 Score 11.8; DB 1; Length 15; Pred. No. 1.4e+02; O; Mismatches 2; Indels
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Pred. No. 1.4e+02;
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Gaps
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US-08-334-847-450

Sequence 450, Application US/08334847 Patent No. 5693532 GENERAL INFORMATION:

APPLICANT:

McSwiggen, James

Kenneth

Draper, Ken Pavco, Pam Woolf, Tod

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RESULT 228
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US-08-334-847-295
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                                                                                                           Matches
                                                                                                                           Query Match
Best Local
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                                                                                                                                                                                                                               TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: McSwig
APPLICANT: Draper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/334,847 FILING DATE: NO. 5693532ember 4, 1994 PRIOR APPLICATION NDATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,3
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: METHOD AND REAGENT FOR TITLE OF INVENTION: INHIBITING RESPIRATORY TITLE OF INVENTION: SYNCYTIAL VIRUS NUMBER OF SEQUENCES: 909

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 633 West Fif
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                        Match 0.8%;
Local Similarity 66.7%;
                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                  TELEFAX:
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                                                                           2235 AGACTATTACAAAAA 2249
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56935;
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1 AAACUAUUACACAAA 15
                                                                                                                                                                                                                     nucleic acid
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633 West Fifth Street
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                                                                                                           Conservative
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Woolf, Tod
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                                                                                                                                                                                        linear
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IBM Compatible
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                                                                                                                        Score 11.8; DB 1;
Pred. No. 1.4e+02;
                                                                                                           Mismatches
                                                                                                                                     Length 15;
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US-08-363-240A-17
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                                                                                                                                                                                                                                                                                                                             Sequence 17, Patent No. 5
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                   TITLE OF
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TITLE OF
TITLE OF
                                                                                                                                                                                                                                            APPLICANT: Couture, Larry
APPLICANT: McSwiggen, James
APPLICANT: Bisgaier, Charles
APPLICANT: Pape, Michael
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                 TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20,
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/334, FILING DATE: No. 5693532ember PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (215) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
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STATE: California
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                                                COUNTRY:
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                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         2310 ATACACTCATCAGAG 2324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11;
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                                      90071
                                                                                                                                                                                                                                                                                                                                7, Application US/08363240A
5705388
                                                                                     Los Angeles
                                                                                                                                                                                INVENTION: METHOD AND REAGENT FOR INVENTION: PREVENTION, INHIBITION OF INVENTION: PROGRESSION AND REGRESSION INVENTION: OF VASCULAR DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INVENTION:
                                                                   California
                                                                                                 Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Suite 4700
                                                                                                                   633 West Fifth Street
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633 West Fifth Street
                                                    U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
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                                                                                                                                  Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%;
73.3%;
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                                                                                                                                                                  1243
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 11.8;
Pred. No. 1.
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                                                                                                                                                                                                  AND REGRESSION
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US-08-363-240A-47/c
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Best Local Similarity 66.7%;
Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION UNMER: 32,327
REFERENCE/DOCKET NUMBER: 210/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 495-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: NOTITLE OF INVENTION: FITTLE OF INVENTION: FITTLE OF INVENTION: COMMERCES:
                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/
FILING DATE: December 23,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Couragen, January McSwiggen, January Bisgaier, Charles Michael
                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: U5/08
FILING DATE: December 23,
PRIOR APPLICATION DATA:
                                                   FILING DATE: ATTORNEY/AGENT INFORMATION:
                                                                                                    FILING DATE: Decembe PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                            COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
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NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 210/096
                                                                                                                                                                                                                                                                                                                                       CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                      STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: sing
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TELEX: 67-3510
                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                    COUNTRY:
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                                                                                                                                                                                                                                                                                                                     California
                                                                                                                                                                                                                                                                                                                                                      633 West Fifth Street Suite 4700
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                                                                                                                                            US/08/363,240A
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Pred. No. 1.4e+02;
                                                                                                                                                                                              DOS 5.0
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TELECOMMUNICATION (213) 407 TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 47
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
TYPE: nucleic single
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                                                 US-08-363-240A-673
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches 13; Conservative
                                                                                                                                          INFORMATION FOR SEQ ID NO:
                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                   NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 210/096
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 1
FILING DATE: December
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
                                                                                                                                            TELEFAX: 12-3510
                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: METHOD AND REAGENT FOR TITLE OF INVENTION: PREVENTION, INHIBITION TITLE OF INVENTION: PROGRESSION AND REGRESSIOTITLE OF INVENTION: OF VASCULAR DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION
                                                                                                                                                         TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 633 West Fifth Street STREET: Suite 4700 CITY: Los Angeles
                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 0.8%;
Local Similarity 86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
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                                                                                          nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Los Angeles
: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    McSwiggen, James
Bisgaier, Charles
Pape, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Couture, Larry
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                                                                                                                                                                                                                                                                                                                                                                                                                                 storage
                                                                                                                                                                                                                                                                                                                              December 23, 1994
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0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PREVENTION, INHIBITION PROGRESSION AND REGREST OF VASCULAR DISEASES
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                                                                                                                                            673:
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Pred. No. 1.
Score 11.8; DB 1;
Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                  DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                1.44 Mb
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AND REGRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.4e+02
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              Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 233
US-08-344-820-9/c
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US-08-317-432A-5
                                                                                         Sequence 9, Application US/08344820 Patent No. 5717085
                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 5, Application US/08317432A Patent No. 5710028
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                                                                        GENERAL INFORMATION:
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APPLICANT: Nurit
 APPLICANT: LYTTLE, NAPPLICANT: KAUVAR, ITTITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/9:
FILING DATE: 27-Jul-92
APPLICATION NUMBER: 08/01
FILING DATE: 1-Jul-93
ATTORNEY/AGENT INFORMATION:
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: MS DOS version 6.2,
OPERATING SYSTEM: Windows version 3.11
SOPTWARE: Word for Windows version 2.0
SOPTWARE: converted to ASCI
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/317,432A
                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 972-3-5625553
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: 1.44 megabyte, 3.5" microdisk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                                                                                                                                       Local Similarity
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TELEPHONE: 9/2-3-5625554
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Friedmam, Mark M.
REGISTRATION NUMBER: 33,883
REFERENCE/DOCKET NUMBER: 12
                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Silver Spring
                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                2631 AAGAAGTTCTTGTTC 2645
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                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
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                                                                                                                                                                                                                                                                                                                                                           ESS: single
linear
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                                                                                                                                                                                                                                                                         Conservative
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LAWRENCE M.

CODON AMIDITES AND METHOD OF USING THEM
TO PRODUCE OLIGONUCLECTIDES AND MUTAGENESIS LIBRARIES
                                                        MATTHEW H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ×
                                                                                                                                                                                                                                                                     0.8%; Score 11.8; DB 1; Length 15; B6.7%; Pred. No. 1.4e+02; tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                       Gaps
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US-08-311-486C-171
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FILING DATE: 23-NOV-1994
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: DROST, PATRICIA M.
REGISTRATION NUMBER: 25,700
REFERENCE/DOCKET NUMBER: 2550
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-1500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                              APPLICANT: Dan T. St
APPLICANT: James MCS
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/344,820
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                     CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
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TELEFAX: 90-4030
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (202) 822-0168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 11.8; DB 1; Length 15; Local Similarity 86.7%; Pred. No. 1.4e+02; les 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 2000 Penn
CITY: Washington
                                                                           OPERATING SYSTEM:
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                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1881 GATGATGAAGATGAT 1895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15 GATGATGATCATGAT 1
                                                                                                                                                                                                                                                                                                                                                              ANT: Kenneth Draper
ANT: Kevin Kisich
ANT: Dan T. Stinchcomb
ANT: James McSwiggen
DF INVENTION: RIBOZYME TREATMENT OF
DF INVENTION: DISEASES OR CONDITIONS
DE INVENTION: RELATED TO LEVELS OF
                                                                                                                                                                          90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20006-1812
                                                                                                                                                                                                                                                                     E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                     Suite 4700
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sean Sullivan
                                                       SYSTEM: IBM P.C. Word Perfect 5.1
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                                                                                           IBM Compatible
September 23, 1994
                                                                                                                                                                                                                                                                                                                              1157
                 US/08/311,486C
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                                                                           DOS 5.0
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US-08-311-486C-172
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NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: (213) 955-0440
TELEFX: 67-3510
JANFORMATION FOR SEQ ID NO: 171:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity Matches '11; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: FITTLE OF 
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
FILING DATE: January 19, 1993
                                                                                                                                                                                                                                                                   SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA: PRIOR APPLICATION DATA: PRIOR APPLICATION DATA:
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FILING DATE: January
APPLICATION NUMBER:
FILING DATE: Decembe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                              FILING DATE: Se
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 633 West Fifth Street
STREET: Suite 4700
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1390 CCAGACTACCTGGAG 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/08311486C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     James McSwiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sean Sullivan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kevin Kisich
Dan T. Stinchcomb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                               IBM Compatible
                                                                                                                                                                               жилькк: US/08/311,486C
September 23, 1994
N: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RIBOZYME TREATMENT OF
DISEASES OR CONDITIONS
RELATED TO LEVELS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1157
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CY 19, 1993
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Pred. No. 1.4e+02;
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G DATE: January 19, CATION NUMBER: 07/9

07/989,849

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                                                                                                                                                                                              PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA: including includi
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APPLICATION UMBER: US/(
FILING DATE: September:
CLASSIFICATION: 435
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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STRANDEDNESS:
TOPOLOGY: lin
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OPERATING SYSTEM:
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TYPE: nucleic acid
STRANDEDNESS: single
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Los Angeles
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.633 West Fifth Street
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James McSwiggen
NVENTION: RIBOZYME TREATMENT OF
NVENTION: DISEASES OR CONDITIONS
NVENTION: RELATED TO LEVELS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Word Perfect 5.1
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Dan T. Stinc
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:

TELEFAX:

(213) 955-0440

STRANDEDNESS:

nucleic acid

ENGTH:

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RESULT 237
US-08-292-620A-242/c
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Matches 11; Conserv
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FILING DATE: December 7, 1992
ATTORNEY AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                  COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. D
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/29
FILING DATE: August 17, 1994
CLASSIFICATION: 435
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TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                PRIOR APPLICATION DATA: PRIOR APPLICATION DATA: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                        ZIP: 90071-2066
COMPUTER READABLE FORM:
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                                                                                                             RIOR APPLICATION DATA: including RIOR APPLICATION DATA: described APPLICATION NUMBER: 08/008,895 FILING DATE: January 19, 193 APPLICATION NUMBER: 07/989,849
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STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kenneth G. Draper
VENTION: RIBOZYME TREATMENT OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sean Sullivan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              James McSwiggen
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73.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DISEASES OR CONDITIONS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RELATED TO LEVELS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2390
                                                                                                                                                                                                                                                   US/08/292,620A
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Pred. No. 1.
                                                                                                                                                                                                                                                                                                       DOS 5.0
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US-08-292-620A-430/c
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Best Local Similarity
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                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including
PRIOR APPLICATION DATA: described
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 193
APPLICATION NUMBER: 07/989,849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Kenneth of TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                             NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                      FILING DATE: December 7, 1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                             Local
                                                                                                                                                                                                                TELEPHONE: (213) 955-0440
                                                                                                                            STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: August
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM CON OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Suite 4700
CITY: Los Angeles
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1650 GCTGGCAGGGGTCTC 1664
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                                                  l Similarity
13; Conserv
                                                                                                                     TH: 15 base pairs nucleic acid NDEDNESS: single LOGY: linear
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INVENTION: RIBOZYME TREATMENT OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INVENTION: DISEASES OR CONDITIONS
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633 West Fifth Street
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Dan T. Stinchcomb
James McSwiggen
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                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SYSTEM: IBM P.C. Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                          August 17, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Stinchcomb
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                                               0.8%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 1.4e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE-1 (I-CAM-1)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US/08/292,620A
                                                                                                                                                                                                                                                                                                                                                                                                  including application described below:
                                                                                                                                                                                                    430:
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Pred. No. 1.
                                                                                                                                                                                                                                                                              208/149
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                                                                    Matches
                                                                                                    Query Match
                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA: including PRIOR APPLICATION DATA: described APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 193
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/14:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION NUMBER: 208/14:
                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
MEDIUM TYPE: STORAGE
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STREET: Suite
STREET: Suite
TTY: Los Angeles
California
                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION: RIBOZYME TREATMENT OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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                                                                    Local Similarity
nes 7; Conser
                                                                                                                                                                  LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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OPERATING SYSTEM:
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                                                                                                                                                         TOPOLOGY:
                                2324 GTGATGTCTGGTCCT 2338
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GUGCUGUAUGGUCCU 15
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633 West Fifth Street
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                                                                                                                                                                                                                                                                              (213) 955-0440
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                                                                      Conservative
                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3: storage
IBM Compatible
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                                                                                 0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    08/008,895
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      including application described below:
                                                                    <u>ه.</u>
                                                                                                                                                                                                                                                                                                                                  208/149
                                                                                 Score 11.8; DB 1;
Pred. No. 1.4e+02;
                                                                    Mismatches
                                                                                                   Length 15;
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RESULT 240 US-08-292-620A-534/c

Sequence 534, Application US/08292620A Patent No. 5837542

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RESULT 241
US-08-292-620A-547
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INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                                 Sequence
                                                                                                GENERAL INFORMAPPLICANT:
APPLICANT:
APPLICANT:
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PRIOR APPLICATION DATA: desc
PRIOR APPLICATION DATA: desc
APPLICATION NUMBER: 08/000
FILING DATE: January 19,
APPLICATION NUMBER: 07/985
FILING DATE: December 7,
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CILAGO DATE: APPLICATION NUMBER: US/OB/OCCIONATE: APPLICATION NUMBER: US/OB/OCCIONATE: APPLIANG DATE: APPLIANG 
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TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                         APPLICANT:
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
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STREET: 633 West Fifth Street
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                                                                                                                                                                                              INFORMATION:
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                                                                                                                                                                                                                    47, Application US/08292620A
5837542
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INVENTION: RIBOZYME TREATMENT OF
INVENTION: DISEASES OR CONDITIONS
INVENTION: RELATED TO LEVELS OF
INVENTION: INTRACELLULAR ADHESION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    California
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                                                                                      Susan Grimm
Dan T. Stinchcomb
James McSwiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (213) 955-0440
                                                             Sean Sullivan
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UMBER: 07/989,849
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3. Draper
RIBOZYME TREATMENT OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A: including application
A: described below:
08/008,895
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Pred. No. 1.
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RESULT 242
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                                                                                                                                                                                                                                      Patent No.
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                             APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
                                                                                                                                                                                                                      GENERAL INFORMATION:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Btorage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
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PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/;
FILING DATE: August 17, 19;
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TITLE OF INVENTION:
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nes 7; Conserv
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STREET: 633 West Fifth Street
STREET: Suite 4700
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James McSwiggen
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Lyon & Lyon
                     ADDRESS:
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August 17, 1994
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633 West Fifth Street

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PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08
FILING DATE: January 1
APPLICATION NUMBER: 07
                                                                                                                                                          TITLE OF INVENTION:
TITLE OF INVENTION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
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COMPUTER READABLE FORM
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                                              CITY: Los Angeles
STATE: California
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REGISTRATION NUMBER: 32,327
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OPERATING SYSTEM:
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nucleic acid
DEDNESS: single
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                  90071-2066
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California
                                                                                                                                                       I: Kenneth G. Draper INVENTION: RIBOZYME TREATMENT OF INVENTION: DISEASES OR CONDITIONS INVENTION: RELATED TO LEVELS OF INVENTION: INTRACELLULAR ADHESION INVENTION: MOLECULE-1 (I-CAM-1)
                                                                                                                                                                                                                                                                                                                                                      , Application US/08292620A
                                                                              Suite 4700
                                                                                             3: Lyon & Lyon
633 West Fifth Street
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James McSwiggen
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Dan T. Stin
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UMBER: 07/989,849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 1.4e+02;
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Best Local :
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                                                                                                                          FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
APPLICATION NUMBER: 07/989,849
AFILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
NAME: WILSON, MARY J. 1955 REGISTRATION NUMBER: 32,955 REFERENCE/DOCKET NUMBER: 55 TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 816-4000
                                                                               APPLICATION NUMBER: US/01
FILING DATE: 07-JUN-1996
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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APPLICANT: PARK, JONG-GU
TITLE OF INVENTION: METHODS OF INHIBITING PHAGOCYTOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
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MEDIUM TYPE:
COMPUTER: IBM
                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20
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1100 NORTH GLEBE ROAD, 8TH FLOOR
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86.7%;
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                                                                                                                                                                   Release #1.0, Version #1.30
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described below:
08/008,895
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                                    555-46
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Pred. No. 1
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RESULT 246
US-08-774-306A-35
; Sequence 35, Application US/08774306A
; Patent No. 5869253
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                       TELEFAX: (703) 01. 2: INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS: 15 base pairs 1.ENGTH: 15 base pairs
                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 13; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No. 5858981 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 55
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 816-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 07-JUN-1996
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: SCHREIBER, ALAN D.
APPLICANT: PARK, JONG-GU
TITLE OF INVENTION: METHODS OF INHIBITING PHAGOCYTOSIS
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER:
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TOPOLOGY: linear
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Y: U.S.A.
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DEDNESS: single
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1100 NORTH GLEBE ROAD, 8TH FLOOR
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RESULT 247
US-08-774-306A-301/c
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Best Local S
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/182,968
FILING DATE: January 13, 1994
APPLICATION NUMBER: 07/882,888
FILING DATE: May 14, 1992
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                    GENERAL
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                                                                                                                                                                                   TITLE OF INVENTION: ITTLE OF INVENTION: TITLE OF INVENTION: NUMBER OF SEQUENCES:
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44
MEDIUM TYPE: storage
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Ly
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CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
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TELEPHONE: (213)
                                                                                                ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
                                                 COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Match 0.8%;
Local Similarity 66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 223/227
                                                                                      STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Com
OPERATING SYSTEM:
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STATE: California
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STREET: 633 West Fifth Street
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                                                                                                                                                                                                                                                                      INFORMATION:
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                                                                                                                                                                                                                                                      Draper, Kenneth G.
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METHOD AND REAGENT FOR INHIBITING HEPATITIS C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IBM P.C. DOS 5.0
                                                                                                                                                                                                      VIRUS REPLICATION
                                                                                                                                                                                                                    METHOD AND REAGENT FOR INHIBITING HEPATITIS C
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Pred. No. 1.
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US-08-774-306A-302/c
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Draper, Kenneth G.
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: INHIBITING HEPATITIS C
TITLE OF INVENTION: VIRUS BEDITARY
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FILING DATE: January 13/
APPLICATION NUMBER: 07/8:
FILING DATE: May 14, 199:
ATTORNEY/AGENT INFORMATION:
                                                                  FILING DATE: May 14, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                          APPLICATION NUMBER: US/08/774,306A FILING DATE: December 26, 1996 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                 SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
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                                                                                                                                                                                                                                         COMPUTER: IBM Com
OPERATING SYSTEM:
                               NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,3
                                                                                                      APPLICATION NUMBER: 08/182,968 FILING DATE: January 13, 1994 APPLICATION NUMBER: 07/882,888
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STATE: California
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   REFERENCE/DOCKET NUMBER: 22:
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633 West Fifth Street
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IUMBER: 07/882,888
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INFORMATION:
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Y 13, 1994
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Pred. No. 1.4e+02;
                223/227
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Best Local Similarity
Matches 13; Conserv
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                            Matches
                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/585
PRILING DATE: January 16, 1996
PRIOR APPLICATION DATA: 60/000,95
PRILING DATE: July 7, 1995
PRILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
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                                                                                                                                                                           TELEFAX: 167-3510
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                                                                                                                                                                                         TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DO:
SOFTWARE: FastSEQ Version 1.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
TOPOLOGY: lin
                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
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1782 TGACAAAGACAAGCC 1796
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o. 5877021
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: California
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633 West Fifth Street
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                              Conservative
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,, 1995
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Pred. No. 1
                                          Score 11.8; DB 1;
Pred. No. 1.4e+02;
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                              Mismatches
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                                                         Length 15;
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RESULT 251
US-08-585-684B-639/c
US-08-585-684B-639/c
Sequence 639, Application US/08585684B
Patent No. 5877021
Patent No. 58770
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; Sequence 638, Application US/08585684B
: Patent No. 5877021
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
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REGISTAN NUMBER: 32,327
REFERENCION NUMBER: 21
REFERENCION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/585,684B
FILING DATE: January 16, 1996
PRIOR APPLICATION DATE:
APPLICATION NUMBER: 60/000,951
FILING DATE: JULY 7, 1995
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: 8torage
MEDIUM TYPE: 8torage
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SOFTWARE: FASTSEQ VERSION 1.5
CURRENT APPLICATION DATA:
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 15 base pairs TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear
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CITY: Los Angeles
STATE: California
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633 West Fifth Street
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US-08-585-684B-640/c
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Best Local Similarity 86.7%;
                                                                                                                                                                                                                                                               APPLICANT: Stinchcomb, Daniel APPLICANT: Jarvis, Thale APPLICANT: McSwiggen, James TITLE OF INVENTION: METHOD ANTITLE OF INVENTION: AND REVER NUMBER OF SEQUENCES: 2751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
               OPERATING SYSTEM: IBM P.C. DO:
SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/000,951
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Wasburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/585,684B FILING DATE: January 16, 1996 PRIOR APPLICATION DATA:
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OPERATING SYSTEM: IBM P.C. DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                     STREET: 633 West F
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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                                          COMPUTER: IBM Com
OPERATING SYSTEM:
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                                                                                                                                   ZIP: 9007:
                                                                                                                                                     COUNTRY:
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STREET: Suite 4700
APPLICATION NUMBER:
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STATE: California
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                                                                                                                                                                                                      Suite 4700
                                                                                                                                                     U.S.A.
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                                                               IBM Compatible
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US/08/585,684B
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 5877021 GENERAL INFORMATION:
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Best Local Similarity 86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM PC.C DOS:
SOFTWARE: FASTESRO VERSION 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,60
FILING DATE: January 16, 1996
FRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/000,951
FILING DATE: JULY 7, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                           TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
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                  SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                     NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 211
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: January PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21
                                                                                                TELEFAX:
                                                                                                  TELEPHONE: (213) 489-16
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
TYPE: _nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1880 AGATGATGAAGATGA 1894
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: California
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                                                                                                                                                              218/078
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; TOPOLOGY: 1: US-08-585-684B-1246
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                                                                                                                                   US-08-585-684B-2267
                                                                                                                                                                                 TELEFAX: (213)
TELEX: 67-3510
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
FURNGTH: 15 base pairs
TENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-585-684B-2267/c
; Sequence 2267, Application US/08585684B
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                                                                  Matches
                                                                                  Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,684B
FILING DATE: January 16, 1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND REA
TITLE OF INVENTION: INDUCTION OF G
TITLE OF INVENTION: AND REVERSAL C
NUMBER OF SEQUENCES: 2751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                    NAME: Warburg, Richard REGISTRATION NUMBER: 32,327 REFERENCE/DOCKET NUMBER: 21 TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/000,951 FILING DATE: July 7, 1995 ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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Local Similarity 60.0%;
hes 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                             Local Similarity
nes 13; Conserv
                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                         TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                               2398 GTGGAGGAACTTTTT 2412
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15 GTGGAGGTACTGTTT 1
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                                                                  Conservative
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                                                                                                                                                     linear
                                                                                0.8%;
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Pred. No. 1.
                                                                                Score 11.8; DB 1; Pred. No. 1.4e+02;
                                                                    Mismatches
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                                                                                                  Length 15;
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                                                                    Indels
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                                                                  Gaps
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RESULT 255 US-09-064-156A-35 ; Sequence 35, Application US/09064156A

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US-09-064-156A-301/c
; Sequence 301, Application US/09064156A
; Patent No. 6132966
                                                                                                                                                                                                                                                                          RESULT 256
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Patent No. 61327.
Paraner
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                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/0
FILING DATE: April 21, 1998
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 234/083
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                         APPLICANT: Draper, Kenneth G.
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: INHIBITING HEPATITIS C
TITLE OF INVENTION: VIRUS REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: May 14, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                       CORRESPONDENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                            STREET: 633 West Fifth Street
STREET: 630 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: January 13, 1994
APPLICATION NUMBER: 07/882,888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/774,306
FILING DATE: December 26, 1996
APPLICATION NUMBER: 08/182,968
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
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STREET: 633 West Fifth Street
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                                                                                                                                                                                                    INFORMATION:
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Los Angeles
California
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VENTION: METHOD AND REAGENT FOR VENTION: INHIBITING HEPATITIS C
VENTION: VIRUS REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                         ADDRESS
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per 26, 1996
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                                                                                                                                                                                                                                                                                                                                                                                                                       Score 11.8;
Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 15;
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RESULT 257
US-09-064-156A-302/c
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Hest Local Similarity 86.7%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 6132966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 302, Application US/09064156A
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APPLICATION NUMBER: 08/774,306
FILING DATE: December 26, 1996
APPLICATION NUMBER: 08/182,968
FILING DATE: January 13, 1994
APPLICATION NUMBER: 07/882,888
FILING DATE: May 14, 1992
ATTORNEY/MERT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                   APPLICATION NUMBER: US/09/064,156A FILING DATE: April 21, 1998 PRIOR APPLICATION DATA: 08/774,306
                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/0
FILING DATE: April 21, 1998
                                                                                                                  COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: WORD PERFECT 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Draper, Kenneth G.
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: INHIBITING HEPATITIS C
TITLE OF INVENTION: VIRUS REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 90071-2066

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                 ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                              COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARB: Word Perfect 5.1
APPLICATION NUMBER: FILING DATE: December APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 234/083
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2436 AATGGATAAGCCAGC 2450
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                      December 26,
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                                                                                                                                                              IBM P.C. DOS 5.0
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Pred. No. 1.4e+02;
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US-09-071-845-242/c
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Best Local Similarity 86.7%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sean Sull APPLICANT: Kenneth G TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                     CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                         SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Dan T. Stinchcom
APPLICANT: James McSwiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: May 14, 199 ATTORNEY/AGENT INFORMATION:
                            FILING DATE: December 7, ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                              APPLICATION NUMBER: US/08/292,620 FILING DATE: August 17, 1994 APPLICATION NUMBER: 08/008,895 FILING DATE: January 19, 1993 APPLICATION NUMBER: 07/989,849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 234/083
                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                           OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (213) 955-0440
NAME: Warburg, Richard REGISTRATION NUMBER: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E OF INVENTION:
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: California
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6132967
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                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                    Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sean Sullivan
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UMBER: 07/882,888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DISEASES OR CONDITIONS RELATED TO LEVELS OF INTRACELLULAR ADHESION
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                                                                                                                                                                                                                                                           IBM P.C. DOS 5.0
                                                                                                                                                                                                         US/09/071,845
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Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-071-845-242
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
TENGTH: 15 base pairs
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Best Local Similarity 86.7%;
Matches 13; Conservative
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                                                                                                                                                                                     APPLICATION NUMBER: US/08/292,620
APPLICATION NUMBER: 08/008,895
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/ACENT INFORMATION:
NAME: WALDURG, Richard J.
REGISTRATION NUMBER: 32,327
TOTAL T
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
AMEDIUM TYPE: STORAGE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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       TELEFAX: 12-
TELEFAX: 67-3510
                                    TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
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VENTION: RIBOZYME TREATMENT OF
VENTION: DISEASES OR CONDITIONS
VENTION: RELATED TO LEVELS OF
VENTION: INTRACELLULAR ADESION
VENTION: INTRACELLULAR ADESION
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linear
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Pred. No. 1.
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SEQUENCE CHARACTERISTICS:

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                                                  US-09-071-845-462
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Query Match
Best Local Similarity
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                                                                                                                                                    TELEFAX: (213) 955-04-
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                 NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: LOB Anna 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TUMBER OF SEQUENCES:
                                                                                                                                                                      TELEPHONE: (213) 955-0440
                                                                  STRANDEDNESS:
TOPOLOGY: 111
                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 90071-2066
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Local Similarity 86.7%;
Les 13; Conservative
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California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       James McSwiggen
Sean Sullivan
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Dan T. Stin
                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IBM Compatible SYSTEM: IBM P.C.
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0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                 US/08/292,620
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                                                                                                                                                        462:
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Score 11.8; DB 1;
Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 11.8; DB 1
Pred. No. 1.4e+02
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            Length 15;
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US-09-071-845-534/c
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                                                                           Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                                                                                                                  NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                  FILING DATE: December 7, 1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                       TELEPAX: 12-
TELEPAX: 67-3510
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                                                   0.8%;
Local Similarity 86.7%;
hes 13; Conservation
                                                                                                                                                                                                                                        TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/292, FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 193
APPLICATION NUMBER: 07/989,849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM COMPONERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                      LENGTH: 15 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Los Angeles
                                                                                                                                        TOPOLOGY:
                          1650 GCTGGCAGGGGTCTC 1664
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VENTION: RIBOZYME TREATMENT OF
VENTION: DISEASES OR CONDITIONS
VENTION: RELATED TO LEVELS OF
VENTION: INTRACELLULAR ADHESION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              James McSwiggen
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                                                                                                                                          linear
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                                                             Pred. No. 1.4
                                                                                         Score 11.8; DB 1;
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                                                        1.4e+02;
2;
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                                                                                         Length 15;
                                                             Indels
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RESULT 263
US-09-071-845-664/c
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TELEPAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 56
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
TYPE: nucleic single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-071-845-547
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Best Local Similarity 45...
Then 7; Conservative
                                                    Sequence 664, Application US/09071845
Patent No. 6132967
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APPLICANT: Susan APPLICANT: Dan T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
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APPLICANT:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REGERENCE/DOCKET NUMBER: 200
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
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APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A. ZIP: 90071-2066
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STATE: California
                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                            2324 GTGATGTCTGGTCCT 2338
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Susan Grimm
Dan T. Stinchcomb
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                                                                                                                                                                                                                                                                                                                                            linear
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                                                                                                                                                                                                                                                                  0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      547:
                                                                                                                                                                                                                                                                  Score 11.8; DB 1; Length 15; Pred. No. 1.4e+02;
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US-09-071-845-730/c
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                                                                                                                                                                   Patent No. 6132967
GENERAL INFORMATION:
                                                                                                                                                                                      Sequence 730, Application US/09071845 Patent No. 6132967
                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.8%;
Best Local Similarity 86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Sean Sull
APPLICANT: Kenneth G
TITLE OF INVENTION:
APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: James MCSWiggen
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: RELATED TO LEVELS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/071,845
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A. ZIP: 90071-2066
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                   1650 GCTGGCAGGGGTCTC 1664
                                                                                                                                                                                                                                                                                                          15 GCTGGTAGAGGTCTC 1
                                                                                                                                                                                                                                                                                                                                                                                      13;
                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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FERNTION: RIBOZYME TREATMENT OF
JENTION: RIBISEASES OR CONDITIONS
FENTION: RELATED TO LEVELS OF
JENTION: INTRACELLULAR ADHESION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       James McSwiggen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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INTRACELLULAR ADHESION

CORRESPONDENCE ADDRESS: TITLE OF INVENTION:

OF SEQUENCES:

MOLECULE-1 (I-CAM-1)

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US-09-038-073-87/c
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base
                                                                                                                                                                                                                                                                              Sequence 87, Application US/09038073 Patent No. 6194150
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,62
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 193
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                                                            APPLICANT: Stinchcomb, Danie
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                 NUMBER OF SEQUENCES:
                                             CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 / Match 0.8%;
Local Similarity 86.7%;
nes 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (213) 489-160
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,3 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                ADDRESSEE:
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READABLE FORM
                                                                              EE: Lyon & Lyon
633 West Fifth Street
Suite 4700
                               U.S.A.
                                                                                                                                                                                                                                               Stinchcomb, Daniel T
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Pred. No. 1.4e+02;
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RESULT 266
US-09-038-073-638/c
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INFORMATION FOR SEQ ID NO: 8
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 638, Application US/09038073 Patent No. 6194150
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                                                                                                                               COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: $1.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/038,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Stinch
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,
REFERENCE/DOCKET NUMBER:
                                                                                 FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Jarvis, Thale APPLICANT: McSwiggen, Jan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE:
MEDIUM TYPE:
COMPUTER: IBN
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Local Similarity 86.7%;
hes 13; Conservative
                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                        CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                                                                                                                                                                              METHOD AND REAGENT FOR THE INDUCTION OF GRAFT TOLERANCE
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              32,327
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 218/078
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Pred. No. 1.
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:

TELEPHONE: TELEFAX: (TELEX: 67-

(213) 489-1600 (213) 955-047

67-3510

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                                           Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. 6194150
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 639, Application US/09038073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                       ATTORNEY AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
TELEDHONE: (213) 489-1600
                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Stinchcomb, Daniel T. APPLICANT: Jarvis, Thale APPLICANT: McSwiggen, James
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADUKESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 0.8%; Score 11.8; DB 1; Length 15; Local Similarity 86.7%; Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                        TELEPHONE: (213) 489-16
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM:
SOFTWARE: FastSE(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Los Angeles
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TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
1880 AGATGATGAAGATGA 1894
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                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     METHOD AND REAGENT FOR THE INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES 2751
                                           0.8%; Score 11.8; DB 1;
86.7%; Pred. No. 1.4e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM P.C.
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                                                                                                                                                                                                                                                                                                                                                                                                   08/585,684
                                                                                                                                                                                                                                                                                                               218/078
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                                                                            Length 15;
                                             0,
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                                          Gaps
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RESULT 268
US-09-038-073-640/c
; Sequence 640, Application US/09038073
"""ant No. 6194150
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                                                                                                                         Sequence 1246, Applic Patent No. 6194150 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 67-3510
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, Jame
TITLE OF INVENTION: METHOL
TITLE OF INVENTION: INDUCT
TITLE OF INVENTION: AND RE
                APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND R.
TITLE OF INVENTION: INDUCTION OF
TITLE OF INVENTION: AND REVERSAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.
SOFTWARE: Fast-SEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/038,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Watburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: sing
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                                                                                                                                                                Application US/09038073
                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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                                                                                                                                                                                                                                                                                                                                               0.8%;
METHOD AND REAGENT FOR THE INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES 2751
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                                                                                                                                                                                                                                                                                                                                                               Length 15;
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CORRESPONDENCE ADDRESS:

ADDRESSEE:

STREET: 633 West Fifth Street

Lyon & Lyon

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TELEX: 67-3510

INFORMATION FOR SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs

TYPE: nucleic acid

TYPE: nucleic single
                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 6194150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 60. Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Stinchcomb, Danie APPLICANT: Jarvis, Thale APPLICANT: McSwiggen, James
             COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FascSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                               CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: 111
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Suite 4700
                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S.A.
                                                                                                                                                                                                                                                         ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 AUGAGGUAUACUUAG 15
                                                                                                                                                  90071
                                                                                                                                                                                                                                    E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                   Suite 4700
                                                                                                                                                                U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                    Stinchcomb, Daniel T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 11.8; DB 1; Length 15; 60.0%; Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                           METHOD AND REAGENT FOR THE INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                           2751
US/09/038,073
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RESULT 271
US-09-158-980-25/c
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Best Local Similarity 86.7%;
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APPLICANT: SCHREIBER, AL
APPLICANT: PARK, JONG-GU
                                                                                                        TELEPHONE: (703) 816-41
TELEFAX: (703) 816-41
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Warburg, NAME: 32,327
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1000
TELEPHONE: (213) 489-1600
TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/657,884
                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
ZIP: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: METHODS OF INHIBITING PHAGOCYTOSIS NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                          NAME: WILSON, MARY J.
REGISTRATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 555-46
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/158,980 FILING DATE: CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: NIXON & VANDERHYE P.C. STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR CITY: ARLINGTON STATE: VIRGINIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single TOPOLOGY: linear
                                 STRANDEDNESS:
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                                                    nucleic acid
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 DNA (genomic)
                                 single
                                                                                                      ) 816-4100
, 816-4100
-- WO: 25:
                                                                                                                                           816-4000
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Pred. No. 1.
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Query Match

Local

Similarity

Length 15;

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; STRANDEDNESS:
; TOPOLOGY: line
; MOLECULE TYPE: I
US-09-158-980-29
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                                                                                                                                                           RESULT 273
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                                                                                       GENERAL INFORMATION:
                                                                                                           Sequence 26, Application US/09081646 Patent No. 6333152
                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                             APPLICANT: Kinzler, Kenneth
APPLICANT: Vogelstein, Bert
APPLICANT: Zhang, Lin
APPLICANT: Zhou, Wei
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IIILE OF INVENTION: Gene Expression Profiles in No. 6333152mal and
IIILE OF INVENTION: Cancer Cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (703)-816-4100 (NFORMATION FOR SEQ ID NO: :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/158,980
                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: WILSON, MARY J.
REGISTRATION NUMBER: 32,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: PARK, JONG-GU
TITLE OF INVENTION: METHODS OF INHIBITING PHAGOCYTOSIS
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                               1334 CTGCATGGTTGACAG 1348
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o. 6242427
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                                                                                                                                                                                                                                                                                                                                                                                          : 15 base pairs nucleic acid
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Y: U.S.A.
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                                                                                                                                                                                                                                                                        Conservative
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86.7%;
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                                                                                                                                                                                                                                                                                     Score 11.8;
Pred. No. 1.
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Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                        Mismatches
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ches 2;
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Sequence 331, Application US/09081646

Patent No. 6333152

GENERAL INFORMATION:

APPLICANT: Kinzler, Kenneth
APPLICANT: Vogelstein, Bert
APPLICANT: Zhang, Lin
APPLICANT: Zhou, Wei
APPLICANT: Zhou, Wei
TITLE OF INVENTION: Gene Expression Profiles in No. 6333152mal and
TITLE OF INVENTION: Cancer Cells
FILE REFERENCE: 01107,74664

CURRENT APPLICATION NUMBER: US/09/081,646

CURRENT FILING DATE: 1998-05-20
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21

NUMBER OF SEQ ID NOS: 871
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; SEQ ID NO 26
; LENGTH; 15
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-081-646-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-081-646-331
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US-09-081-646-331
                                                                                                                                                                                                                                                          Sequence 418, Application US/09081646 Patent No. 6333152 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 331 LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.8%;
Best Local Similarity 86.7%;
Matches 13; Conservative
TITLE OF INVENTION: Gene Expression Profiles
TITLE OF INVENTION: Cancer Cells
FILE REFERENCE: 01107.74664
CURRENT APPLICATION NUMBER: US/09/081,646
CURRENT FILING DATE: 1998-05-20
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEO ID NOS: 871
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CURRENT APPLICATION NUMBER: US/09/081,646
CURRENT FILING DATE: 1998-05-20
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ ID NOS: 871
                                                                                                                                                                        APPLICANT: Kinzler, Kenneth
APPLICANT: Vogelstein, Bert
APPLICANT: Zhang, Lin
APPLICANT: Zhou, Wei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 3.0
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86.7%;
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Pred. No. 1.4e+02;
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Pred. No. 1.
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APPLICANT: Bertino, Joseph R.
APPLICANT: Banerjee, Debabrata
APPLICANT: Tong, Youzhi
APPLICANT: Tong, Youzhi
APPLICANT: Liu-Chen, Xinyue
TITLE OF INVENTION: Mutants of Thymidylate Synthase and
FILE REFERENCE: D5978
FULE REFERENCE: D5978
CURRENT APPLICATION NUMBER: US/09/367,007C
CURRENT FILING DATE: 1999-10-15
PRIOR APPLICATION NUMBER: PCT/US98/02145
PRIOR FILING DATE: 1998-01-03
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-081-646-445
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                                                                                                           SEQ ID NO 1
LENGTH: 15
TYPE: DNA
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Best Local Similarity 86.7%;
Matches 13; Conservative
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APPLICANT: Vogelstein, Bert
APPLICANT: Zhang, Lin
APPLICANT: Zhang, Lin
APPLICANT: Zhang, Lin
APPLICANT: Zhong, Wei
TITLE OF INVENTION: Gene Expression Profiles in No. 6333152mal and
TITLE OF INVENTION: Cancer Cells
FILE REFERENCE: 01107.74664
CURRENT APPLICATION NUMBER: US/09/081,646
CURRENT FILING DATE: 1998-05-20
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 445, Application US/09081646
Patent No. 6333152
                                                                                                                                                                              NUMBER OF SEQ ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 871
FEATURE: FEATURE: OTHER INFORMATION: Nucleotide sequence in the 5' coding region of OTHER INFORMATION: human recombinants cDNA of thymidylate synthase OTHER INFORMATION: (TS) gene in pET-17(bhTS) vector.
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ORGANISM: Homo sapiens
                                                                                  ORGANISM: artificial sequence
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; LENGTH: 15
; TYPE: DNA
; ORGANISM: human
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US-09-475-947A-69/c
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US-08-826-134-32/c
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Matches 13
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APPLICANT: Peles, Blior
TITLE OF INVENTION: CASPR/190, A FUNCTIONAL LIGAND FOR RPTP-BETA AND THE
TITLE OF INVENTION: AXONAL CELL RECOGNITION MOLECULE CONTACTIN
FILE REFERENCE: 7683-111
CURRENT TELLING DATE: 1997-03-26
CURRENT FILING DATE: 1997-03-26
EARLIER APPLICATION NUMBER: 60/614,199
EARLIER FILING DATE: 1996-03-27
NUMBER: FILING DATE: 1996-03-27
                                                                                                                                                                                                                                                                                                         FILE REFERENCE: UTSD0667
CURRENT APPLICATION NUMBER: US/09/475,947A
CURRENT FILING DATE: 1999-12-31
NUMBER OF SEQ ID NOS: 346
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 69
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Best Local Similarity 86.7%;
Matches 13; Conservative
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Garner, Harold R.
APPLICANT: Wren, Jonathan D.
APPLICANT: Minna, John D.
TITLE OF INVENTION: Polymorphic Repeats in Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE: OTHER INFORMATION: Description of Artificial Sequence: primer
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Pred. No. 1.4e+02;
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Pred. No. 1.4e+02;
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37 12.8 0.9 17 1 US-09-817-879-4457 38 12.8 0.9 17 1 US-10-392-970-29 39 12.8 0.9 17 1 US-10-392-970-29 40 12.8 0.9 17 1 US-10-209-787-1750 41 12.8 0.9 17 1 US-10-209-787-3314 42 12.8 0.9 17 1 US-10-209-787-3315 43 12.8 0.9 17 1 US-10-209-787-3315 44 12.8 0.9 17 1 US-10-209-787-3315 45 12.8 0.9 17 1 US-10-060-830-700 46 12.8 0.9 17 1 US-10-060-830-700 47 12.8 0.9 17 1 US-10-060-756A-144 48 12.8 0.9 17 1 US-10-060-756A-144 49 12.8 0.9 17 1 US-10-060-756A-145 49 12.8 0.9 17 1 US-10-060-756A-1675 50 12.8 0.9 17 1 US-10-060-756A-1677 50 12.8 0.9 17 1 US-10-060-998-588 51 12.8 0.9 17 1 US-10-060-998-589 52 12.8 0.9 17 1 US-10-163-552-677	219 12.8 0.9 17 1 US-09-745-237A-366 221 12.8 0.9 17 1 US-09-745-237A-660 221 12.8 0.9 17 1 US-09-745-237A-660 221 12.8 0.9 17 1 US-09-745-237A-1169 22 12.8 0.9 17 1 US-09-72-818-283 224 12.8 0.9 17 1 US-10-238-700-661 222 12.8 0.9 17 1 US-10-238-700-324 225 12.8 0.9 17 1 US-10-238-700-324 226 12.8 0.9 17 1 US-10-238-700-3544 226 12.8 0.9 17 1 US-10-238-700-3544 227 12.8 0.9 17 1 US-10-238-700-3544 229 12.8 0.9 17 1 US-10-238-700-121 239 12.8 0.9 17 1 US-10-61-201-121 239 12.8 0.9 17 1 US-10-61-201-121 239 12.8 0.9 17 1 US-09-817-879-483 231 12.8 0.9 17 1 US-09-817-879-483 233 12.8 0.9 17 1 US-09-817-879-483 235 12.8 0.9 17 1 US-09-817-879-1421 235 12.8 0.9 17 1 US-09-817-879-1421 235 12.8 0.9 17 1 US-09-817-879-1325 236 12.8 0.9 17 1 US-09-817-879-3135 236 12.8 0.9 17 1 US-09-	182 12.8 0.9 17 1 US-09-818-875-315-1760 184 12.8 0.9 17 1 US-09-818-875-31760 185 12.8 0.9 17 1 US-09-818-875-315-1760 185 12.8 0.9 17 1 US-09-818-875-315-1760 185 12.8 0.9 17 1 US-09-818-875-315-1760 185 12.8 0.9 17 1 US-09-818-875-315-186 12.8 0.9 17 1 US-09-780-533A-2092 17 1 US-09-780-533A-2092 17 1 US-09-780-533A-2092 17 1 US-09-780-533A-2103 191 12.8 0.9 17 1 US-09-848-754A-184 192 12.8 0.9 17 1 US-09-848-754A-184 194 12.8 0.9 17 1 US-09-848-754A-184 197 12.8 0.9 17 1 US-09-848-754A-186 199 12.8 0.9 17 1 US-09-848-754A-3109 12.8 0.9 17 1 US-09-930-423-1169 12.8 0.9 17 1 US-09-780-164-525 12.8 0.9 17 1 US-09-827-395A-611 12.8 0.9 17 1 US-09-740-332-99 12.8 0.9 17 1 US-09-740-332-99 12.8 0.9 17 1 US-09-740-332-653 12.8 0.9 17 1 US-09-740-332-653 12.8 0.9 17 1 US-09-740-332-1321 12.8 0.9 17 1 US-09-740-332-3322 13.8 0.9 17 1 US-09-740-33	0 12.8 0.9 17 1 US-09-864-785-664
37 12.8 0.9 17 1 US-09-817-879-4457 Sequence 2 12.8 0.9 17 1 US-10-392-970-29 Sequence 2 12.8 0.9 17 1 US-10-299-787-1759 Sequence 2 12.8 0.9 17 1 US-10-209-787-1759 Sequence 1 12.8 0.9 17 1 US-10-209-787-1760 Sequence 1 12.8 0.9 17 1 US-10-209-787-3314 Sequence 1 12.8 0.9 17 1 US-10-209-787-3315 Sequence 1 12.8 0.9 17 1 US-10-209-787-3315 Sequence 2 12.8 0.9 17 1 US-10-600-830-700 Sequence 2 12.8 0.9 17 1 US-10-600-830-700 Sequence 2 12.8 0.9 17 1 US-10-600-756A-144 Sequence 2 12.8 0.9 17 1 US-10-600-756A-146 Sequence 2 12.8 0.9 17 1 US-10-600-756A-1675 Sequence 2 12.8 0.9 17 1 US-10-600-756A-1675 Sequence 2 12.8 0.9 17 1 US-10-600-998-588 Sequence 2 12.8 0.9 17 1 US-10-600-998-589 Sequence 2 12.8 0.9 1	219 12.8 0.9 17 1 US-09-745-237A-346 220 12.8 0.9 17 1 US-09-745-237A-366 221 12.8 0.9 17 1 US-09-745-237A-660 Sequence 3 221 12.8 0.9 17 1 US-09-745-237A-1660 Sequence 6 222 12.8 0.9 17 1 US-09-745-237A-1169 Sequence 6 223 12.8 0.9 17 1 US-09-728-188-283 Sequence 2 224 12.8 0.9 17 1 US-10-238-700-661 Sequence 2 225 12.8 0.9 17 1 US-10-238-700-3248 Sequence 2 226 12.8 0.9 17 1 US-10-238-700-3544 Sequence 3 227 12.8 0.9 17 1 US-10-238-700-3544 Sequence 3 228 12.8 0.9 17 1 US-10-238-701-121 Sequence 1 230 12.8 0.9 17 1 US-10-61-201-121 Sequence 1 231 12.8 0.9 17 1 US-09-817-879-99 232 12.8 0.9 17 1 US-09-817-879-481 Sequence 3 233 12.8 0.9 17 1 US-09-817-879-483 Sequence 4 233 12.8 0.9 17 1 US-09-817-879-1421 Sequence 4 234 12.8 0.9 17 1 US-09-817-879-1421 Sequence 3 235 12.8 0.9 17 1 US-09-817-879-313 Sequence 3 236 12.8 0.9 17 1 US-09-817-879-310 Sequence 3 237 128 0.9 17 1 US-09-817-879-310 Sequence 3 238 128 0.9 17 1 US-09-817-879-310 Sequence 3 239 129 129 129 129 129 129 129 129 129 12	182 12.8 0.9 17 1 US-09-818-875-1759 Sequence 185 12.8 0.9 17 1 US-09-818-875-1759 Sequence 185 12.8 0.9 17 1 US-09-818-875-1759 Sequence 185 12.8 0.9 17 1 US-09-818-875-3315 Sequence 186 12.8 0.9 17 1 US-09-818-875-3315 Sequence 187 12.8 0.9 17 1 US-09-818-875-3315 Sequence 188 12.8 0.9 17 1 US-09-818-875-3315 Sequence 189 12.8 0.9 17 1 US-09-780-533A-2103 Sequence 191 12.8 0.9 17 1 US-09-780-533A-2103 Sequence 191 12.8 0.9 17 1 US-09-848-754A-184 Sequence 192 12.8 0.9 17 1 US-09-848-754A-184 Sequence 193 12.8 0.9 17 1 US-09-848-754A-184 Sequence 194 12.8 0.9 17 1 US-09-848-754A-184 Sequence 195 12.8 0.9 17 1 US-09-848-754A-184 Sequence 196 12.8 0.9 17 1 US-09-848-754A-184 Sequence 197 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 198 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 207 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 208 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 208 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 209 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 200 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 200 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 200 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 201 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 202 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 203 12.8 0.9 17 1 US-09-848-754A-3109 Sequence 204 12.8 0.9 17 1 US-09-876-474-679 Sequence 205 12.8 0.9 17 1 US-09-878-164-594 Sequence 206 12.8 0.9 17 1 US-09-878-164-594 Sequence 207 12.8 0.9 17 1 US-09-878-38A-691 Sequence 208 12.8 0.9 17 1 US-09-878-38A-691 Sequence 208 12.8 0.9 17 1 US-09-873-35A-691 Sequence 208 12.8 0.9 17 1 US-09-873-35A-691 Sequence 208 12.8 0.9 17 1 US-09-740-332-481 Sequence 208 12.8 0.9 17 1 US-09-740-332-395 Sequence 208 12.8 0.9 17 1 US-09-7	0 12.8 0.9 17 1 US-09-864-785-664 Sequence 6

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RESULT 1
US-09-073-881-4/c
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 Sequence 4, Application US/09073881
Patent No. US20020045251A1
GENERAL INFORMATION:
COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.5 inch, 1.

COMPUTER: Compaq Presario 4540

OPERATING SYSTEM: Windows 95

SOFTWARE: Word Perfect 8.0

CURRENT APPLICATION NUMBER: US/09/073,881

FILING DATE:

CLASSIFICATION:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/073,881

FILING DATE: 07-MAY-1997

ATTORNEY/AGENT INFORMATION:

NAME: Alan J. Howarth

REGISTRATION NUMBER: 36,553

REFERENCE/DOCKET NUMBER: 74903.CIP

TELEPHONE: (801)566-6633

TELEPHONE: (801)566-663
 CORRESPONDENCE ADDRESS:
ADDRESSEE: Thorpe, No. US20020045251Alth & Western, STREET: P.O. Box 1219
 APPLICANT: Rao, Mahendra S.
APPLICANT: Mujtaba, Tahmina
AP
 NUMBER OF SEQUENCES:
 CITY: Sandy
STATE: Utah
 USA
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 36,553
ER: T4903.CIP
 US-10-163-552-935
 -09-837-992-29
-09-816-460C-26
 ALIGNMENTS
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174, App
9628, App
157, Appl
57, Appl
911, App
50, Appl
192, App
2272, Ap
 935, App
364, App
365, App
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1457, Ap
1545, Ap
2834, Ap
4426, Ap
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 ; OTHER INFORMATION: Antisense Oligonucleotide US-09-954-556-62
 RESULT 3
US-09-954-556-62/c
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 US-09-954-556-61
 RESULT 2
US-09-954-556-61/c
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 US-09-073-881-4
 CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 62
LENGTH: 20
 GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPREFILE REFERENCE: RTS-0250
 GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR
FILE REFERENCE: RTS-0250
FILE REFERENCE: RTS-0250
 Query Match
Best Local Similarity
Matches 20; Conserv
 Sequence 62, Application US/09954556 Publication No. US20030078219A1
 Query Match
Best Local Similarity
Matches 20; Conserv
 SEQ ID NO 61
 Sequence 61, Application US/09954556 Publication No. US20030078219A1
 Best Local Similarity Matches 20; Conserv
 Query Match
 CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
 INFORMATION FOR SEQ ID NO:
 ORGANISM: Artificial Sequence
 TYPE: DNA
 OTHER INFORMATION: Antisense Oligonucleotide
 TYPE: DNA ORGANISM: Artificial Sequence
 FEATURE:
 FEATURE:
 LENGTH: 20
 SEQUENCE CHARACTERISTICS:
 TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
 LENGTH:
 1335
 1318 GGGATATCCTTTCACTCTGC 1337
 1317
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 20
 TGCATGGTTGACAGTTCTGC 1354
 TGGGATATCCTTTCACTCTG 1336
 TGCATGGTTGACAGTTCTGC 1
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 1.4%; Score 20;
100.0%; Pred. No.
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 Length 20;
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US-09-954-556-65/c
; Sequence 65, Application US/09954556
; Publication No. US20030078219A1
; GENERAL INFORMATION:
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 RESULT 6
 US-09-954-556-64/c
 US-09-954-556-64
 RESULT 5
 ; OTHER INFORMATION: Antisense Oligonucleotide US-09-954-556-63
 US-09-954-556-63/c
 SEQ ID NO 64
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
 Sequence 64, Application US/09954556 Publication No. US20030078219A1. GENERAL INFORMATION:
 SEQ ID NO 63
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
 Sequence 63, Application US/099 Publication No. US20030078219A1 GENERAL INFORMATION:
 Query Match
Best Local 9
 Matches
 Matches
 Query Match
 APPLICANT:
 APPLICANT: SUBAN M. Freier
APPLICANT: SCOTT COOPER
TITLE OF INVENTION: ANVISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRE
FILE REFERENCE: RTS-0250
CURRENT FLICATION NUMBER: US/09/954,556
CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FLILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
 TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPREFILE REFERENCE: RTS-0250
CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
 APPLICANT: Brett P. Monia
APPLICANT: Susan M. Frei
APPLICANT: Scott Cooper
 APPLICANT: Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRE
 APPLICANT: Brett P. Monia
 OTHER INFORMATION: Antisense Oligonucleotide
 FEATURE:
 FEATURE:
 Local
 Local
 1411 ATTTACTGCATAGGGGTCTT 1430
 1421 TAGGGGTCTTCTTAATCGCC 1440
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 20 TAGGGGTCTTCTTAATCGCC
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 Similarity
 Susan M. Freier
Scott Cooper
 Susan M. Freier
Scott Cooper
 Application US/09954556
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 1.4%; Score 20;
100.0%; Pred. No.
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 Score 20; Pred. No.
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 Length 20;
 Length 20;
 Indels
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; OTHER INFORMATION: Antisense Oligonucleotide
US-09-954-556-67
 US-09-954-556-67/c
 ; OTHER INFORMATION: Antisense Oligonucleotide US-09-954-556-66
 US-09-954-556-66/c
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 US-09-954-556-65
 CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 67
LENGTH: 20
TYPE: DNA
 GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Susan M. Frei
 CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 66
LENGTH: 20
 FILE REFERENCE: RTS-0250
CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 65
LENGTH: 20
 Sequence 67, Application US/09954556 Publication No. US20030078219A1
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Best Local Similarity
 GENERAL INFORMATION:
 Sequence 66, Application US/09954556 Publication No. US20030078219A1
 Matches
 Query Match
 APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
TITLE OF INVENTION: ANTISENSE MODULATION OF
 APPLICANT: Brett P. Monia
APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0250
 FILE REFERENCE: RTS-0250
 ORGANISM: Artificial Sequence FEATURE:
 FEATURE:
 ORGANISM: Artificial Sequence
 TYPE: DNA
 OTHER INFORMATION: Antisense Oligonucleotide
 ORGANISM: Artificial Sequence FEATURE:
 TYPE: DNA
 Local Similarity es 20; Conserv
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 1479 CACGACCAAGAAGCCAGACT 1498
 20
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 CACGACCAAGAAGCCAGACT 1
 Conservative
 Conservative
 1.4%;
 100.0%;
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 Score 20;
Pred. No.
 Score 20;
Pred. No.
 Mismatches
 Mismatches
 DB 1;
24;
 B
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 FIBROBLAST GROWTH FACTOR RECEPTOR
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 Length 20
 Length 20;
 Indels
 Indels
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 0
 Gaps
 Gaps
 0
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 2 EXPRE
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Query Match

Score 20;

DB

1;

Length

20;

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APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF FIB
FILE REFERENCE: RTS-0250
CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 69
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE: VICTUARITY M. Artigenes Oligonicleotide
 APPLICANT: Susan M. Freier

APPLICANT: Susan M. Freier

FILE REFERENCE: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRINTER REPORT APPLICATION NUMBER: US/09/954,556

CURRENT FILING DATE: 2001-09-14

NUMBER OF SEQ ID NOS: 108

LENGTH: 20

TYPE: DNS
 RESULT 11
US-09-954-556-70/c
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 RESULT 10
US-09-954-556-69/c
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 RESULT 9
US-09-954-556-68/c
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 US-09-954-556-68
Sequence 70, Application US/09954556 Publication No. US20030078219A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
Matches 20; Conserv
 Query Match 1.4
Best Local Similarity 100
Matches 20; Conservative
 Sequence 69, Application US/09954556 Publication No. US20030078219A1
 Best Local Similarity Matches 20; Conserv
 GENERAL INFORMATION:
 OTHER INFORMATION: Antisense Oligonucleotide
.09-954-556-69
 TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
 OTHER INFORMATION: Antisense Oligonucleotide
 1489
 2028 GTACTCCTATGACATTAACC 2047
 2023 ATGGAGTACTCCTATGACAT 2042
 20
 20
 20
 AAGCCAGACTTCAGCAGCCA 1
 AAGCCAGACTTCAGCAGCCA 1508
 Conservative
 Conservative
 1.4%; Score 20; DB 1;
100.0%; Pred. No. 24;
Live 0; Mismatches
 1.4%; Score 20; DB 1;
100.0%; Pred. No. 24;
 100.0%; Pred. No.
 0; Mismatches
 <u>.</u>
 Mismatches
 FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPR
 <u>.</u>
 0
 Length 20
 Length 20
 Indels
 Indels
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 Gaps
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 US-09-954-556-72/c
 RESULT 12
US-09-954-556-71/c
 RESULT 13
 US-09-954-556-70
 Matches
 Query Match
 Query Match
 Matches
 LENGTH: 20
TYPE: DNA
 ORGANISM: Artificial Sequence
 FEATURE:
 20
 20
```

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; OTHER INFORMATION: Antisense Oligonucleotide US-09-954-556-71
Sequence 72, Application US/09954556
Publication No. US20030078219A1
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRE FILE REFERENCE: RT9-0250
CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 72
LENGTH: 20
 CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 71
LENGTH: 20
 GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
 NUMBER OF SEQ ID NOS: 108
SEQ ID NO 70
LENGTH: 20
 Sequence 71, Application US/09954556 Publication No. US20030078219A1
 TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPREFILE REFERENCE: RTS-0250
 APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR
FILE REFERENCE: RTS-0250
FILE REFERENCE: RTS-0250
 CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
 TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
Artificial Sequence
FEATURE:
Artificial Sequence
 APPLICANT: Brett P. Monia
 TYPE: DNA
ORGANISM: Artificial Sequence
 OTHER INFORMATION: Antisense Oligonucleotide
 Local Similarity hes 20; Conserv
 Local Similarity
mes 20; Conserv
 2387 CAGGGATTCCCCGTGGAGGAA 2406
 2032 TCCTATGACATTAACCGTGT 2051
 CAGGGATTCCCGTGGAGGAA 1
 1.4%; Score 20;
larity 100.0%; Pred. No.
Conservative 0; Mismatc
 Conservative (
 1.4%; Score 20;
100.0%; Pred. No.
 0,
 Mismatches .
 Mismatches
 DB 1;
. 24;
 DB 1;
 0
 0
 Length 20;
 Length 20;
 Indels
 0
 0
 Gaps
 Gaps
 0;
 0
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RESULT 16
 US-09-954-556-74/c
 ; OTHER INFORMATION: Antisense Oligonucleotide
US-09-954-556-73
 US-09-954-556-73/c
 US-09-954-556-74
 RESULT 15
 SEQ ID NO 74
LENGTH: 20
TYPE: DNA
 APPLICANT: Brett P. Monia
APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
APPLICANT: SCOTT COOPER
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRE
FILE REFERENCE: RTS-0250
CURRENT APPLICATION UNMBER: US/09/954,556
CURRENT APPLICATION UNMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 73
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
 Sequence 73, Applicat Publication No. US200 GENERAL INFORMATION:
 Matches
 Best Local Similarity Matches 20; Conserv
 Best Local Similarity Matches 20; Conserv
 Matches
 GENERAL INFORMATION:
 Sequence 74, Application US/09954556 Publication No. US20030078219A1
 Query Match
 Query Match
 Query Match
 APPLICANT: SUSAN M. Freier
APPLICANT: Scott Cooper
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRE
 NUMBER OF SEQ ID NOS: 108
 CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
 OTHER INFORMATION: Antisense Oligonucleotide
 OTHER INFORMATION: Antisense Oligonucleotide
 ORGANISM: Artificial Sequence
 FEATURE:
 FEATURE:
 Local Similarity
 2562 CACAACCAATGAGGAATACT 2581
 2557
 2536 GAAGACTIGGATCGAATICI 2555
 20;
 20
 20
 20 GAAGACTTGGATCGAATTCT
 ACTCTCACAACCAATGAGGA 2576
 CACAACCAATGAGGAATACT 1
 Application US/09954556
 Conservative
 Conservative 0;
 Conservative
 1.4%; 5...
100.0%; Pred. No.
10e 0; Mismatches
 100.0%;
 1.4%; Score 20;
L00.0%; Pred. No.
 1.4%; Score 20;
00.0%; Pred. No.
 <u>,,</u>
 Mismatches
 Mismatches
 DB 1;
24;
 DB
24;
 DB 24;
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 0,
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 0;
 Length 20;
 Length 20;
 Length 20
 Indels
 Indels
 0
 <u>,,</u>
 0,
 Gaps
 Gaps
 Gaps
 0
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 0
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 RESULT 17
US-09-954-556-96/c
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 US-09-954-556-98/c
 ; OTHER INFORMATION: Antisense Oligonucleotide US-09-954-556-96
 ; OTHER INFORMATION: Antisense Oligonucleotide US-09-954-556-75
 US-09-954-556-75/c
 FILE REPERENCE: RTS-0250
CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 75
LENGTH: 20
 Sequence 75, Application US/09954556 Publication No. US20030078219A1 GENERAL INFORMATION:
 Sequence 98, Application US/09954556 Publication No. US20030078219A1 GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
APPLICANT: SCOTT COOPER
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRE
FILE REFERENCE: RTS-0250
CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 98
 SEQ ID NO 96
LENGTH: 20
 Matches
 Query Match
 GENERAL INFORMATION:
 Sequence 96, Application US/09954556 Publication No. US20030078219A1
 Matches
 Query Match
 CURRENT APPLICATION NUMBER: US/09/954,556
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 108
 APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0250
 APPLICANT: Brett P. Monia
APPLICANT: Susan M. Frei
APPLICANT: Scott Cooper
 APPLICANT: Brett P. Monia
APPLICANT: Susan M. Freier
APPLICANT: Scott Cooper
TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRE
 TYPE: DNA ORGANISM: Artificial Sequence
 TYPE: DNA
ORGANISM: Artificial Sequence
 FEATURE:
 FEATURE:
 Local Similarity nes 20; Conserv
 Local Similarity 100.0%; Pred. No. les 20; Conservative
 2591 GCCAACCTCTCGAACAGTAT 2610
 2701 CCTCAGTATCCACACATAAA 2720
 20 GCCAACCTCTCGAACAGTAT 1
 20 CCTCAGTATCCACACATAAA 1
 Conservative
 100.0%;
 1.4%; Score 20;
00.0%; Pred. No.
 0;
 Mismatches
 Mismatches
 DB 1;
 DB 1;
 FIBROBLAST GROWTH FACTOR RECEPTOR 2 EXPRE
 <u>.</u>.
 °,
 Length 20
 Length 20;
 Indels
 Indels
 0;
 0,
 Gaps
 Gaps
 0
 0
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APPLICANT: Gill, Parkesh
APPLICANT: MASOOd, Rizwan
APPLICANT: MASOOd, Rizwan
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: VEGF OLIGONUCLEOTIDES
FILE REFERENCE: 21327-701CON2
CURRENT APPLICATION NUMBER: US/09/805,761
CURRENT FILING DATE: 2001-03-13
PRIOR APPLICATION NUMBER: PCT/US01/00019
PRIOR PILING DATE: 2001-01-19
PRIOR PILING DATE: 2001-01-19
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 US-09-805-761-43
 RESULT 20
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 US-09-805-761-42
 US-09-954-556-98
 Sequence 43, Application US/09805761 Patent No. US20020165174A1 GENERAL INFORMATION:
 SEQ ID NO 42
LENGTH: 23
 Query Match
 OTHER INFORMATION: VEGFR-1 gene specific primers for RT-PCR-09-805-761-42
 GENERAL INFORMATION:
APPLICANT: Gill, Pa
APPLICANT: Masood,
 Query Match
 Sequence 42, Appl
Patent No. US2002
 TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR ANTISENSE TITLE OF INVENTION: VEGF OLIGONUCLEOTIDES FILE REFERENCE: 21327-701CON2
 PRIOR APPLICATION NUMBER: US 09/487,023
PRIOR FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: US 09/016,541
PRIOR FILING DATE: 2000-11-24
 NUMBER OF SEQ ID NOS: 64.
SOFTWARE: FastSEQ for Windows Version 4.0
 CURRENT APPLICATION NUMBER: US/09/805,761
 PRIOR FILING DATE: 2001-01-19
 PRIOR APPLICATION NUMBER: PCT/US01/00019
 TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
 OTHER INFORMATION: Antisense Oligonucleotide
 FEATURE:
 TYPE: DNA
 ORGANISM: Artificial Sequence
 Local Similarity 90.9%;
 Local
APPLICATION NUMBER: US 09/487,023 FILING DATE: 2000-01-19
 2098 CAGCTGGCCAGAGGCATGGAGT 2119
 FILING DATE: 1997-01-31
 APPLICATION NUMBER: US 60/037,004
 APPLICATION NUMBER: US 09/016,541 FILING DATE: 1998-01-30
 2470 TACATGATGAGGGACTG 2489
 20
 Similarity
 CAAGTGGCCAGAGGCATGGAGT 22
 Application US/09805761
 Masood, Rizwan
 Conservative
 Conservative
 0165174A1
 Parkesh
 2001-03-1
 1.4%; Score 20; DB 1; 100.0%; Pred. No. 24;
 0
 Score 18.8;
Pred. No. 4
 Mismatches
 Mismatches
 47;
 DB 1; Length 23;
 FOR ANTISENSE
 <u>ب</u>
 0;
 Length 20;
 Indels
 0
 0
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RESULT 22
US-10-094-466-79
; Sequence 79, Application US/10094466
; Publication No. US20030203363A1
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 RESULT 21
US-09-953-047-5/c
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 US-09-953-047-5
 US-09-805-761-43
 GENERAL INFORMATION
 Sequence 5, App. Publication No.
 Matches
 SEQ ID NO 5
 GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Jacqueline Wy:
 SOFTWARE: FastSEQ for Windows Version SEQ ID NO 43
 Best Local Similarity
 Query Match
 Matches
 Best Local Similarity
Matches 20; Conservat
 Query Match
 CURRENT APPLICATION NUMBER: US/10/094,466
CURRENT FILING DATE: 2002-03-07
PRIOR APPLICATION NUMBER: 60/274,281
PRIOR FILING DATE: 2001-03-08
PRIOR APPLICATION NUMBER: 60/288,148
PRIOR FILING DATE: 2001-05-02
PRIOR FILING DATE: 2001-05-02
PRIOR APPLICATION NUMBER: 60/274,849
 APPLICANT: Spytek et al.
TITLE OF INVENTION: NOVEL HUMAN PROTEINS,
TITLE OF INVENTION: AND METHODS OF USING
TITLE OF INVENTION: THE SAME
FILE REFERENCE: 21402-290D
 CURRENT APPLICATION NUMBER: US/09/953,047
CURRENT FILING DATE: 2001-09-10
 TITLE OF INVENTION: ANTISENSE MODULATION OF FIBROBLAST GROWTH FACTOR RECEPTOR 3 EXPRE
 PRIOR APPLICATION NUMBER: US 09/016,541
PRIOR FILING DATE: 2000-11-24
PRIOR APPLICATION NUMBER: US 09/016,541
PRIOR FILING DATE: 1998-01-30
PRIOR APPLICATION NUMBER: US 60/037,004
 NUMBER OF SEQ ID NOS: 95
 PRIOR FILING DATE: 1997-01-31
NUMBER OF SEQ ID NOS: 64
 ORGANISM: Artificial Sequence FEATURE:
 TYPE: DNA
 LENGTH: 23
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
 OTHER INFORMATION: PCR Primer
 OTHER INFORMATION: VEGFR-1 gene specific primers
 LENGTH: 21
APPLICATION NUMBER: 60/275,235
 1822 AAGATGTTGAAAGATGATGCC 1842
 2098 CAGCTGGCCAGAGGCATGGAGT 2119
 21 AAGATGCTGAAAGACGATGCC 1
 5, Application US/09953047
 Conservative
 Conservative
 US20030087854A1
 1.3%;
 -03-09
 90.9%;
 Score 17.8;
Pred. No. 5
 Score 18.8;
Pred. No. 47
 Mismatches
 Mismatches
 POLYNUCLEOTIDES ENCODING THEM
 ВB
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 1;
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 for RT-PCR
 Length 23;
 Indels
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FILING DATE: 2001-03-12

APPLICATION NUMBER: 60/338,375

2001-12-04

LING DATE:

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 ; OTHER INFORMATION: Synthetic Sequence US-09-774-809-62
 RESULT 23
US-09-774-809-62/c
 US-10-094-466-79
 Sequence 62, Appropriate Publication No.
 Matches
 Best Local Similarity
 Query Match
 Matches
 Best Local Similarity
 Query Match
 CURRENT APPLICATION NUMBER: US/09/774,809
CURRENT FILING DATE: 2001-01-31
PRIOR APPLICATION NUMBER: 09/396,902
 PRIOR APPLICATION NUMBER: 09/130,616
PRIOR FILING DATE: 1998-08-07
PRIOR APPLICATION NUMBER: 08/910,629
PRIOR FILING DATE: 1997-08-03
 APPLICANT: Gaarde, William A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
FILE REFERENCE: ISPH-0412
 NUMBER OF SEQ ID NOS: 165
 PRIOR FILING DATE: 1999-09-15
 APPLICANT: McKay, Robert A.
APPLICANT: Dean, Nicholas
 SOFTWARE: Patin 2.1
EQ ID NO 79
 Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 114
 APPLICANT:
 ORGANISM: Artificial Sequence FEATURE:
 TYPE: DNA
 OTHER INFORMATION: Description of Artificial Sequence: Forward Primer
 TYPE: DNA
ORGANISM: Artificial Sequence
 FEATURE:
 ENGTH: 22
 ENGTH:
 ID NO 62
 APPLICATION NUMBER: 60/277,338 FILING DATE: 2001-03-20
 FILING DATE: 2001-10-31
APPLICATION NUMBER: 60/275,601
FILING DATE: 2001-03-13
APPLICATION NUMBER: 60/276,000
 1886 TGAAGATGATTGGGAAACACAA 1907
 FILING DATE: 2001-03-14
 APPLICATION NUMBER: 60/335,302
 2642 GTTCTTCAGGAGATGAT 2658
 FILING DATE:
 APPLICATION NUMBER: 60/275,579
 INFORMATION:
 19; Conservative
17
 20
 Monia,
Monia,
Tro, Pam
 TGAACATGTTTGGAAAACACAA 22
 Dean, Nicholas M.
 Application US/09774809
 Conservative
 US20030004120A1
 Brett
 2001-03-13
 1.2%;
 1.2%;
 0;
 Score 17;
Pred. No.
 0; Mismatches
 Score 17.2;
Pred. No. 71;
 Mismatches
 DB 1;
63;
 DB
 1;
 Length 20;
 Length 22
 Indels
 0,
 0
 Gaps
 Gaps
 0,
 0
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RESULT 24

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В
 US-09-989-339-51
 US-09-073-881-21
 US-09-073-881-21
 Sequence 21, Application US/09073881 Patent No. US20020045251A1 GENERAL INFORMATION:
 APPLICANT: Falco,
APPLICANT: Famodu
APPLICANT: Rafals
APPLICANT: Ramake
 GENERAL INFORMATION:
 Sequence 51, Ap
Publication No.
 Matches
 Query Match
NUMBER OF SEQ ID NOS: 55
SOFTWARE: Microsoft Office 97
 PRIOR APPLICATION NUMBER: 08/703,829 PRIOR FILING DATE: 1996-08-27
 CURRENT APPLICATION NUMBER: US/09/989,339
CURRENT FILING DATE: 2002-05-31
 APPLICANT: Thorpe, Catherine TITLE OF INVENTION: PLANT METHIONINE SYNTHASE GENE AND METHODS TITLE OF INVENTION: METHICNINE CONTENT OF THE SEEDS OF PLANTS FILE REFERENCE: BB-1067-B
 TELEFAX: (801)566-0750 INFORMATION FOR SEQ ID NO:
 PPLICANT:
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/8:
FILING DATE: 07-MAY-1997
ATTORNEY/AGENT INFORMATION:
 SEQUENCE CHARACTERISTICS:
 SOFTWARE: Word Perfect 8.0 CURRENT APPLICATION DATA:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
 APPLICANT: Rao, Mahendra S.
APPLICANT: Mujtaba, Tahmina
TITLE OF INVENTION: A Common Neural Progenitor for the CNS and
 TELECOMMUNICATION INFORMATION:
 CORRESPONDENCE ADDRESS:
 NUMBER OF SEQUENCES:
 APPLICATION NUMBER: FILING DATE: CLASSIFICATION:
 COMPUTER: Compaq Presario 4540
OPERATING SYSTEM: Windows 95
 Local Similarity
les 17; Conserv
 STRANDEDNESS:
TOPOLOGY: lir
 REGISTRATION NUMBER: 36 REFERENCE/DOCKET NUMBER:
 TELEPHONE:
 NAME: Alan J. Howarth
 ZIP: 84091-1219
 COUNTRY:
 CITY: Sandy
 STREET:
 LENGTH:
 ADDRESSEE:
 1873 GAGATGGAGATGATGAA 1889
 σ
 nucleic acid
 GAGATGAGATGAA 22
 Famodu, Layo
 Utah
 Rafalski, Jan A.
Ramaker, Michael
 Tarczynski, Mitchell C.
 Application US/09989339
 E: Thorpe, No. P.O. Box 1219
 USA
 Conservative
 linear
 US20030088886A1
 (801)566-6633
 Saverio Carl
 single
 100.0%; ---
 No.
 08/852,744
 US/09/073,881
 36,553
 21:
 US20020045251A1th & Western, L.L.P.
 3.5 inch,
 Score 17;
Pred. No.
 T4903.CIP
 Mismatches
 1.44 Mb storage
 DB 1; Length 22; 75;
 0
 Indels
 0
 FOR INCREASING THE
 PNS
 Gaps
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 US-09-735-995-39/c
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 RESULT 27
 ; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-953-318-72
 ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic oligonucleotide US-09-989-339-51
 US-09-953-318-72/c
 SEQ ID NO 72
LENGTH: 20
TYPE: DNA
 Query Match
Best Local S
Matches 17
 SEQ ID NO 39
LENGTH: 20
TYPE: DNA
ORGANISM: Homo sapiens
 Sequence 39, Application US/09735995 Patent No. US20010034024A1
 GENERAL INFORMATION:
 Sequence 72, Application Us/USS Publication No. US20030105036A1
 Query Match
Best Local Similarity
Matches 18; Conserv
 Matches
 Query Match
Best Local Similarity
 APPLICANT: Splawski, Igor
TITLE OF INVENTION: MUTATIONS IN AND GENOMIC STRUCTURE
TITLE OF INVENTION: SYNDROME GENE
FILE REFERENCE: 2223-136
CURRENT APPLICATION NUMBER: US/09/735,995
CURRENT FILING DATE: 2000-12-14
 APPLICANT: C. Frank Bennett
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: RTS-0232
 PRIOR APPLICATION NUMBER: 09/226,012
PRIOR FILING DATE: 1999-01-06
NUMBER OF SEQ ID NOS: 116
SOFTWARE: Patentin Ver. 2.0
 CURRENT APPLICATION NUMBER: US/09/953,318
CURRENT FILING DATE: 2001-09-13
NUMBER OF SEQ ID NOS: 154
 APPLICANT: Keating, Mark T.
APPLICANT: Splawski, Igor
 ORGANISM: Artificial Sequence
 TYPE: DNA
 ORGANISM: Artificial Sequence FEATURE:
2556 CACTCTCACAACCAATGAG 2574
 2098 CAGCTGGCCAGAGGCATGGA 2117
 2686 TACGAACCATGCCTTCCTCA 2705
 INFORMATION:
 20 CAAGTGGCCAGAGGCATGGA 1
 18;
 Similarity
 TACTAACCATGGCTTCCTCA 20
 Application US/09953318
 Conservative
 Conservative
 1.2%;
 1.2%;
 1.1%;
 ; Score 15.8; D; Pred. No. 91; 0; Mismatches
 <u>.</u>
 0;
 Score 16.8;
Pred. No. 67;
 Score 16.8;
Pred. No. 67;
 Mismatches
 Mismatches
 OF VASCULAR ENDOTHELIAL GROWTH FACTOR RECEPT
 DВ
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 DB 1;
 1; Length 20;
 2
 Length 20;
 Length 20;
 Indels
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 밁
 US-09-848-754A-3493
 RESULT 29
US-09-848-754A-3493
 뭉
 á
 US-09-733-294A-61
 RESULT 28
US-09-733-294A-61
 밁
 GENERAL INFORMATION:
APPLICANT: Brett P.
APPLICANT: William
 Sequence 61, App
Patent No. US200
 Matches
 Query Match
 Matches
 Query Match
 APPLICANT:
 TYPE: DNA
 Local Similarity
 12;
 19
```

```
Sequence 3493, Application US/09848754A
Publication No. US20030073207A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Tr
TITLE OF INVENTION: Levels of Epidermal Grov
FILE REFERENCE: MBHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
 RESULT 30

US-10-055-728-142/c

; Sequence 142, Application US/10055728

; Publication No. US20030170720A1

; GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0
SEQ ID NO 3493
LENGTH: 17
 CURRENT APPLICATION NUMBER: US/09/733,294A
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 09/572,423
PRIOR FILING DATE: 2000-05-16
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 61
LENGTH: 20
 APPLICANT: van der Kuyl, Antoinette C. APPLICANT: Cornelissen, Marion
 APPLICANT: Cornelissen, Marion
TITLE OF INVENTION: MEANS AND METHODS FOR TREATMENT EVALUATION
 APPLICANT: Susan M. Freier
APPLICANT: Edward V. Wancewicz
TITLE OF INVENTION: ANTISENSE MODULATION OF TERT EXPRESSION
FILE REFERENCE: 5244US (REN/P55190US00)
 FILE REFERENCE: ISPH-0527
 TYPE: RNA
ORGANISM: Homo sapiens
 ORGANISM: Artificial Sequence FEATURE:
 OTHER INFORMATION: Antisense Oligonucleotide
 1.1%;
Local Similarity 89.5%;
1es 17; Conservativo
 2318 ATCAGAGTGATGTCTGG 2334
 2020 GGGATGGAGTACTCCTATG 2038
 ACCAGAGUGAUGUCUGG
 GGGATGGACTATTCCTATG 19
 William Gaarde
 Application US/09733294A
 Conservative
 20045588A1
 Monia
 Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
 1.1%;
 Levels of Epidermal Growth
 17
 0
 Score 15.8;
Pred. No. 91
 Score 15.4;
Pred. No. 76;
 Mismatches
 Mismatches
 91;
 DB
 BB
 ۲.
 Factor Receptors
 Length 17;
 Length 20;
 Indels
 0
 <u>,</u>
 Gaps
 Gaps
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 US-08-459-455-89/c
 ; OTHER INFORMATION: 3'TAG019GENE-2 US-10-055-728-142
 Sequence 89, Application US/084
Publication No. US20030124105A1
GENERAL INFORMATION:
 SEQ ID NO 142
 Matches
 Query Match
 Best Local
 PRIOR APPLICATION NUMBER: EP 0120373.2
PRIOR FILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: EP 01200228.3
PRIOR FILING DATE: 2001-01-23
NUMBER OF SEQ ID NOS: 156
 SOFTWARE: PatentIn version 3.1
 CURRENT APPLICATION NUMBER: US/10/055,728
CURRENT FILING DATE: 2002-04-19
 PRIOR APPLICATION NUMBER: 60/325,722
PRIOR FILING DATE: 2001-09-28
 FILING DATE: 24-JUN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence B.
REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0600
TELECOMMUNICATION INFORMATION:
 TYPE: DNA ORGANISM: Artificial Sequence
 NFORMATION FOR SEQ ID NO:
 APPLICANT: Yuan, Junying
APPLICANT: Miura, Masayuki
TITLE OF INVENTION: Programmed Cell Death
NUMBER OF SEQUENCES: 95
 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/258,287
FILING DATE: 10-UN-1994
SEQUENCE CHARACTERISTICS:
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/080,850
 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
 SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
 CORRESPONDENCE ADDRESS:
 ENGTH: 18
 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
 TELEPHONE: (202) 3
TELEFAX: (202) 371
TELEX: 248636 SSK
 APPLICATION NUMBER: US 08/368,704 FILING DATE: 4-JAN-1995
 FILING DATE:
 STATE: D.C.
 CLASSIFICATION:
 APPLICATION NUMBER:
 CITY: Washington
 STREET:
 ADDRESSEE:
 2413 AAGCTGCTGAAGGAAGG 2429
 18
 20005
 Similarity
 AAGCTGCTGAAGGAGGG 2
 Application US/08459455
 E: Sterne, Kessler, Goldstein & Fox 1100 New York Avenue, Suite 600
 USA
 Conservative
 2-JUN-1995
 371-2540
 2002-04-19
 371-2600
 94.1%;
 US/08/459,455
 ; Score 15.4; D; Pred. No. 85; 0; Mismatches
 DB 1; Length 18;
 Genes
 and Proteins
 0
 Gaps
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 US-08-459-455-89
 OTHER INFORMATION: Sense Primer US-09-849-901-4
 RESULT 33
US-09-849-901-4/c
 ; OTHER INFORMATION: Antisense Oligonucleotide US-09-967-655-59
 US-09-967-655-59/c
 1.1%;
Best Local Similarity 94.1%;
Matches 16; Conservative
 GENERAL INFORMATION:
APPLICANT: Wagener, Christoph
TITLE OF INVENTION: Method for Detecting Mutated Alleles
FILE REFERENCE: 4121-111 CIP
CURRENT APPLICATION NUMBER: US/09/849,901
CURRENT FILING DATE: 2001-05-03
 GENERAL INFORMATION:
APPLICANT: C. Frank
APPLICANT: Andrew
 SEQ ID NO 59
LENGTH: 20
 Sequence 59, Application US/09967655 Publication No. US20030092649A1
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 4
Query Match
Best Local Similarity
 Sequence 4, Application US/09849901 Patent No. US20020045227A1
 Matches
 Query Match
Best Local Similarity
 CURRENT APPLICATION NUMBER: US/09/967,655
CURRENT FILING DATE: 2001-09-28
NUMBER OF SEQ ID NOS: 95
 TITLE OF INVENTION: ANTISENSE MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACTOR RECEPT TITLE OF INVENTION: EXPRESSION FILE REFERENCE: RTS-0227
 PRIOR FILING DATE: 1998-03-04
NUMBER OF SEQ ID NOS: 10
 PRIOR APPLICATION NUMBER: DE19708758.2
PRIOR FILING DATE: 1997-03-04
PRIOR APPLICATION NUMBER: PCT/DE98/00676
 PRIOR APPLICATION NUMBER: US 09/390,545
PRIOR FILING DATE: 1999-09-02
 ORGANISM: Artificial Sequence
 TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
 FEATURE
 TYPE: DNA
 ENGTH:
 TOPOLOGY: both
 STRANDEDNESS:
 LENGTH:
 2110 GGCATGGAGTACTTGGC 2126
 1878 GGAGATGATGAAGATGA 1894
 18 GGCATGGAGTTCTTGGC
 20
 16;
 nucleic acid
DEDNESS: both
 GGAGTTGATGAAGATGA 4
 Andrew T.
 20 base pairs
 Frank Bennett
 Conservative
 Watt
1.1%;
 1.1%;
 N
 <u>,</u>
 0;
 Score 15.4;
Pred. No. 1e
 Score 15.4; DB 1;
Pred. No. 1e+02;
Score 15.2; DB 1;
Pred. No. 1.1e+02;
 Mismatches
 Mismatches
 1e+02;
 DB 1;
 Length 20
 Length 20;
 Length 20
 Indels
 Indels
 0
 0
 0
 0
```

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TITLE OF INVENTION: ANTISENSE MODULATION OF TRANSFORMING GROWTH FACTOR-BETA 3 EXPRESS, FILE REFERENCE: RTS-0257
; CURRENT APPLICATION UNMERS: US/09/906,158
; CURRENT FILING DATE: 2001-07-14
NUMBER OF SEQ ID NOS: 168
SEQ ID NO 67
LENGTH
 US-09-906-158-67/c
 RESULT 35
 ₽
 ; NAME/KEY: modified base
; LOCATION: (3)..(4)
; OTHER INFORMATION: gm
US-09-944-036-19
 밁
 US-09-906-158-67
 US-09-944-036-19
 RESULT 34
 Matches
Query Match
Best Local Similarity
 Matches
 Query Match
Best Local
 SEQ ID NO 19
 Sequence 19, Application US/09944036 Patent No. US20020055095A1
 APPLICANT: BRENTANO, Steven T.

APPLICANT: BABOLA, Odile
APPLICANT: TRAN, Nathalie
APPLICANT: VERNET, GUENCES OF HIV-1 SEQUENCES FOR DETECTION
TITLE OF INVENTION: SEQUENCES ASSOCIATED WITH DRUG-RESISTANCE MUTA
 NUMBER OF SEQ ID NOS: 70
SOFTWARE: PatentIn Ver. 2.1
 FILE REFERENCE: GP114-02.UT
CURRENT APPLICATION NUMBER: US/09/944,036
CURRENT FILING DATE: 2001-08-31
 PRIOR APPLICATION NUMBER: US 60/229,790 PRIOR FILING DATE: 2000-09-01
 LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
 OTHER INFORMATION: Description of Artifici OTHER INFORMATION: Oligonucleotide primer OTHER INFORMATION: sequence NAME/KBY: modified base LOCATION: (1)...(2)
OTHER INFORMATION: 2'-O-methyladenosine
 APPLICANT: YANG,
 FEATURE:
OTHER INFORMATION:
 TYPE: DNA
ORGANISM: Artificial Sequence
 FEATURE:
 ENGTH: 20
 2422 AAGGAAGGACACAGAATGGA 2441
 2424 GGAAGGACACAGAATGGATA 2443
 INFORMATION:
 20
 17;
 _
 17;
 Similarity
 AAGGAAGGACACCAAATGAA 20
 GGAAGGAGACAGAGTGGAAA 1
 Conservative
 Conservative
 Yeasing Y.
 Antisense Oligonucleotide
 2001-08-3
 1.1%;
1.1%;
 Score 15.2; DI
Pred. No. 1.1e-
0; Mismatches
 0
 <u>,</u>
Score 15.2; DB 1;
Pred. No. 1.1e+02;
 of Artificial Sequence:
 Mismatches
 1.1e+02;
 for
 DB 1;
 Protease target
 ω
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 Length 20;
 Length 20;
 Indels
 Indels
 0
 0
 Gaps
 Gaps
 0
 0
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RESULT 38
US-10-007-010-59/c
US-10-007-010-59/c
; Sequence 59, Application US/10007010
; Publication No. US20030125275A1
; GENERAL INFORMATION:
 밁
 밁
 US-10-400-670-4
 RESULT 37
 S
 ; OTHER INFORMATION: Antisense Oligonucleotide US-09-953-318-78
 RESULT 36
US-09-953-318-78/c
 밁
 US-10:400-670-4/c
 SEQ ID NO 4
LENGTH: 20
TYPE: DNA
 Sequence 4, Application US/10400670
Publication No. US20030215854A1
GENERAL INFORMATION:
APPLICANT: CLAUSEN, PETER A.
TITLE OF INVENTION: DETECTION OF DI
 CURRENT APPLICATION NUMBER: US/09/953,318
CURRENT FILING DATE: 2001-09-13
NUMBER OF SEQ ID NOS: 154
SEQ ID NO 78
LENGTH: 20
 Query Match
Best Local Similarity
Matches 17; Conser
 Query Match
Best Local
 Sequence 78, Application US/09953318 Publication No. US20030105036A1 GENERAL INFORMATION:
 Matches
 Matches
 CURRENT APPLICATION NUMBER: US/10/400,670 CURRENT FILING DATE: 2003-06-28 NUMBER OF SEQ ID NOS: 11 SOFTWARE: Patentin Ver. 2.1
 TITLE OF INVENTION: ANTISENSE MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACTOR RECEPT TITLE OF INVENTION: EXPRESSION
 FILE REFERENCE: 39147-0013
 APPLICANT: C. Frank Bennett
APPLICANT: Andrew T. Watt
 FILE REFERENCE: RTS-0232
 TYPE: DNA
ORGANISM: Artificial Sequence
 OTHER INFORMATION: OTHER INFORMATION:
 ORGANISM: Artificial Sequence FEATURE:
 FEATURE:
 Local Similarity 85.0%;
mes 17; Conservative
 1718 CACTGGGCAAGCCCCTGGGA 1737
 2275 AAGTGGATGGCTCCAGAAGC 2294
 2423 AGGAAGGACACAGAATGGAT 2442
 20
 20
 20
 17;
 CACTGGGGAATCCCCTTGGA 1
 AAATGGATGGCTCCCGAATC 1
 Conservative
 Conservative
 Description of Artificial hairpin oligonucleotide
 DETECTION OF DNA-BINDING
 1.1%;
 0
 0
 0
 Score 15.2; D
Pred. No. 1.1e
0; Mismatches
 Score 15.2; DB 1;
Pred. No. 1.1e+02;
 Mismatches
 Mismatches
 1.1e+02;
 DB 1;
 PROTEINS
 Sequence: Synthetic
 ω
 Length 20,
 Length 20
 Indels
 Indels
 Indels
 0,
 0;
 0
 Gaps
 Gaps
 Gaps
 0
 0;
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US-10-440-850-753
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 5
 ; TYPE: RNA; Homo sapiens; ORGANISM: Homo sapiens US-10-277-494-59
 ; OTHER INFORMATION: US-10-007-010-59
 US-10-277-494-59
 RESULT 39
 SEQ ID NO 59
LENGTH: 20
TYPB: DNA
ORGANISM: Artificial Sequence
 SEQ ID NO 59
LENGTH: 15
 Sequence 753, Appropriation No.
 Sequence 59, Ap
Publication No.
 Matches
 Query Match
Best Local Similarity
 Query Match
Best Local :
 GENERAL INFORMATION:
 Matches
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Induction of Graft Tolerance
TITLE OF INVENTION: Immune Responses
FILE REFERENCE: 250/130 (MBHB00-900-A)
CURRENT APPLICATION NUMBER: US/10/440,850
CURRENT FILING DATE: 2003-05-19
PRIOR APPLICATION NUMBER: US/09/550,012
PRIOR PELICATION NUMBER: US/09/550,012
 SENERAL INFORMATION:
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or conditions Related To Leve.
TITLE OF INVENTION: Epidermal Growth Factor Receptors
TILE REFERENCE: MBHB00-958-K (400/064)
CURRENT APPLICATION UNMBER: US/10/277,494
CURRENT FILING DATE: 2002-10-21
 APPLICANT: Ribozyme Pharmaceuticals,
APPLICANT: Stinchcomb, Dan
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, Jim
 PRIOR APPLICATION NUMBER: US 60/000,951 PRIOR FILING DATE: 1995-07-07
 NUMBER OF SEQ ID NOS: 446
SOFTWARE: PatentIn version 3.0
 CURRENT APPLICATION NUMBER: US/10/007,010 CURRENT FILING DATE: 2001-12-04
 APPLICANT: Kenneth W. Dobie TITLE OF INVENTION: ANTISENS FILE REFERENCE: RTS-0345
 NUMBER OF SEQ ID NOS: 87
 FEATURE:
APPLICATION NUMBER: US 09/038,073
 2320 CAGAGTGATGTCTGG 2334
 2275 AAGTGGATGGCTCCAGAAGC 2294
 20 AAGTGGACAGCTCCTGAAGC 1
 11;
 1 CAGAGUGAUGUCUGG 15
 Similarity
 Application US/10277494
 Application US/10440850
 Conservative
 Conservative
 US20030207837A1
 US20030186909A1
 Antisense Oligonucleotide
 ANTISENSE MODULATION OF HCK EXPRESSION
 1.1%;
 1.1%;
 0;
 Score 15;
Pred. No.
 Score 15.2; DB 1;
Pred. No. 1.1e+02;
 Mismatches
 68;
 Length 15
 Length 20;
 0;
 0
 Gaps
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 RESULT 42
US-09-848-754A-2907
 US-10-440-850-754
 US-10-440-850-754
 US-10-440-850-753
Sequence 2907, Application US/09848754A
Publication No. US20030073207A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate.
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MSHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
SOFTWARE: PatentIn version 3.0
SEQ ID NO 2907
LENGTH: 17
 PRIOR FILING DATE: 1998-03-11
NUMBER OF SEQ ID NOS: 2285
SOFTWARE: PatentIn version 3.0
SEQ ID NO 753
 1.1%;
Best Local Similarity 73.3%;
Matches 11; Conservative
 SEQ ID NO 754
 GENERAL INFORMATION:
 Sequence 754, Application US/10440850 Publication No. US20030207837A1
 Matches
 Query Match
 Best Local Similarity
 PRIOR FILING DATE: 1998-03-11
NUMBER OF SEQ ID NOS: 2285
SOFTWARE: PatentIn version 3.0
 PRIOR APPLICATION NUMBER: US 08/585,684
PRIOR FILING DATE: 1996-01-12
PRIOR APPLICATION NUMBER: US 60/000,951
PRIOR FILING DATE: 1995-07-07
PRIOR APPLICATION NUMBER: US 09/038,073
 FILE REFERENCE: 250/130 (MBHB00-900-A) CURRENT APPLICATION NUMBER: US/10/440,850 CURRENT FILING DATE: 2003-05-19 PRIOR APPLICATION NUMBER: US/09/650,012 PRIOR FILING DATE: 2000-08-28
 APPLICANT: Stinchcomb, Dan
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Induction of Graft Tolerance and Rever
TITLE OF INVENTION: Immune Responses
FILE REFERENCE: 250/130 (MBHB00-900-8)
FILE REFERENCE: 250/130 (MBHB00-900-8)
 APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Stinchcomb, Dan
 LENGTH: 15
TYPE: RNA
ORGANISM: Mus musculus
 LENGTH: 15
TYPE: RNA
ORGANISM: Mus musculus
 1374 GGAGATTACAGCTTC 1388
 1373 AGGAGATTACAGCTT 1387
 11;
 H
 GGAGAUUACAGCUUC
 AGGAGAUUACAGCUU 15
 Conservative
 1.18;
 15
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 Score 15; DE
Pred. No. 68;
 Score 15;
 Mismatches
 Mismatches
 No.
 68;
 DB
 ۲,
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 Length 15;
 Length 15;
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 Gaps
 Gaps
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US-10-251-117-712
; Sequence 712, Application US/10251117
; Cariantion No. US20030170891A1
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 ; ORGANISM: Homo sapiens
US-09-848-754A-2907
 US-10-251-117-1019/c
 PRIOR APPLICATION NUMBER: US 10/163,552
PRIOR FILING DATE: 2002-06-06
PRIOR PELLCATION NUMBER: US 60/358,580
PRIOR FILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US 09/916,466
PRIOR APPLICATION NUMBER: US 60/296,249
PRIOR FILING DATE: 2001-66-06
 Sequence 1019, Application US/10251117 Publication No. US20030170891A1
 Best Local Similarity 73.3 Matches 11; Conservative
 Matches
 Best Local
 Query Match
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 712
 Query Match
 GENERAL INFORMATION:
 GENERAL INFORMATION:
 CURRENT APPLICATION NUMBER: US/10/251,117
CURRENT FILING DATE: 2003-02-24
PRIOR APPLICATION NUMBER: US 60/393,924
PRIOR FILING DATE: 2002-07-03
PRIOR APPLICATION NUMBER: US 10/163,552
PRIOR FILING DATE: 2002-06-06
 APPLICANT: McSwiggen, James
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Epidermal Growth Factor
TITLE OF INVENTION: Gene Expression Using Short Interfering RNA
FILE REFERENCE: 900/042 (MBHB02-468-A)
 PRIOR APPLICATION NUMBER: US 60/393,924 PRIOR FILING DATE: 2002-07-03
 FILE REFERENCE: 900/042 (MBHB02-468-A)
CURRENT APPLICATION NUMBER: US/10/251,117
CURRENT FILING DATE: 2003-02-24
 NUMBER OF SEQ ID NOS: 1213
 TITLE OF INVENTION: RNA Interference Mediated Inhibition of Epidermal Growth Factor TITLE OF INVENTION: Gene Expression Using Short Interfering RNA
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
 OTHER INFORMATION: Description of Artificial Sequence:
 TYPE: RNA
 FEATURE:
 ORGANISM: Artificial Sequence
APPLICATION NUMBER: US 60/358,580 FILING DATE: 2002-02-20
 2320 CAGAGTGATGTCTGG 2334
 2320 CAGAGTGATGTCTGG 2334
 N
 Similarity
 CAGAGUGAUGUCUGG 16
 CAGAGUGAUGUCUGG 16
 Conservative
 1.1%;
 2003-02-24
 73.3%;
 Score 15; DB 1;
Pred. No. 1.1e+02
 4
 Score
Pred.
 Mismatches
 Mismatches
 No.
 DB 1;
 Length 19
 Length 17;
 Indels
 Target sequence/siNA sense
 0
 °,
 Gaps
 Gaps
 0
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Query Match
Best Local Similarity
Matches 15; Conserva
 RESULT 46
US-10-086-156-49
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 US-09-908-410-14
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 US-09-908-410-14
 US-10-251-117-1019
 SEQ ID NO 14
LENGTH: 20
TYPE: DNA
 APPLICANT: Hogan, Kirk J.
APPLICANT: Brunson, David B.
APPLICANT: Roberts, Monica C.
APPLICANT: Mickelson, James R
TITLE OF INVENTION: Assay For
FILE REFERENCE: 960296.98148
 GENERAL INFORMATION:
APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: POLYNUCLEOTIDE ENCODING
TITLE OF INVENTION: K+betaM4 and K+betaM5
 Sequence 49, Application US/10086156 Publication No. US20030054989A1
 Sequence 14, Application US/09908410 Publication No. US20030104382A1
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 1019
 GENERAL INFORMATION:
 Matches
 Query Match
CURRENT APPLICATION NUMBER: US/10/086,156
CURRENT FILLNG DATE: 2002-02-28
PRIOR APPLICATION NUMBER: US 60/272,190
PRIOR FILING DATE: 2001-02-28
PRIOR APPLICATION NUMBER: US 60/274,258
PRIOR APPLICATION NUMBER: US 60/274,258
PRIOR FILLNG DATE: 2001-03-07
NUMBER OF SEQ ID NOS: 98
 Best Local Similarity
Matches 15; Conserva
 CURRENT APPLICATION NUMBER: US/0
CURRENT FILING DATE: 2001-07-18
NUMBER OF SEQ ID NOS: 20
 FILE REFERENCE: D0115NP
 NUMBER OF SEQ ID NOS: 1213
 PRIOR APPLICATION NUMBER: US 60/296,249 PRIOR FILING DATE: 2001-06-06
 SOFTWARE: PatentIn Ver. 2.1
 PEATURE:
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence:oligonucleotide
NAME/KEY: misc feature
LOCATION: (1)...(20)
OTHER INFORMATION: Human RYR bases 6686-6667.
 ORGANISM: Artificial Sequence FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: sina antisense region
 ORGANISM: Artificial Sequence
 TYPE: RNA
 ENGTH: 19
 2529
 2320 CAGAGTGATGTCTGG 2334
 18
 տ
 GTTGGTAGAAGACTT 2543
 Brunson, David B.
Roberts, Monica C.
 GTTGGTAGAAGACTT 19
 Mickelson, James R
 Conservative
 Conservative
 1.1%;
 1.1%;
 For Propensity For Canine Malignant Hyperthermia
 US/09/908,410
 0;
 <u>,</u>
 Score 15;
Pred. No.
 Score 15;
Pred. No.
 Mismatches
 Mismatches
 B
 1.1e+02;
 DB 1;
 TWO NOVEL HUMAN POTASSIUM CHANNEL BETA-SU
 Length 19;
 Length 20
 Indels
 Indels
 <u>,</u>
 0;
 Gaps
 Gaps
 0
 0
```

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GENERAL INFORMATION:

APPLICANT: Serup, Palle
APPLICANT: Heimberg, Harry
APPLICANT: Gradwohl Gerard
TITLE OF INVENTION: Methods For Generating Insulin-Secreting
TITLE OF INVENTION: Cells Suitable for Transplantation
FILE REFERENCE: 6246.200-US
CURRENT APPLICATION UNMEER: US/10/090,011
CURRENT APPLICATION NUMBER: US 60/271,474
PRIOR APPLICATION NUMBER: US 60/271,474
PRIOR APPLICATION NUMBER: US 60/271,474
PRIOR FILING DATE: 2001-02-26
NUMBER OF SEQ ID NOS: 70
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 38
LENGTH: 20
TYPE: DNA
ORGANISM: Homo Sapien
US-10-090-011-38
 Sequence 30, Application US/10309175
Publication No. US20030148347A1
GENERAL INFORMATION:
APPLICANT: Georg A. HOLLANDER
TITLE OF INVENTION: NUCLEAR PROTEIN
FILE REFERENCE: 117-427 / N84006
CURRENT APPLICATION NUMBER: US/10/309,175
CURRENT FILING DATE: 2002-12-04
PRIOR APPLICATION NUMBER: US 60/336,176
PRIOR FILING DATE: 2001-12-06
NUMBER OF SEQ ID NOS: 32
SOFTWARE: MS WORT
LENGTH: 18
 RESULT 48
US-10-309-175-30/c
 片
 S
 RESULT 47
US-10-090-011-38/c
 밁
 ş
 ; LENGTH: 20
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-086-156-49
 ; PEATURE:
; NAME/KEY: misc feature
; OTHER INFORMATION: Description of Artificial Sequence: Primer
US-10-309-175-30
 SEQ ID NO 49
LENGTH: 20
 Sequence 38, Application US/10090011 Publication No. US20030082810A1
 Matches
 Query Match
 Matches 15; Conservative
 Query Match
Best Local Similarity
 Query Match
 Matches
 ORGANISM: Artificial sequence
 TYPE: DNA
 Local Similarity 100.0%;
 Match 1.1%;
Local Similarity 88.9%;
 1807 GTCACCGTGGCCGTG 1821
 1691 AATGGGAGTTTCCAA 1705
 15; Conservative
 16 GTCACCGTGGCCGTG 2
16; Conservative
 AATGGGAGTTTCCAA 20
 1.1%; Score 15;
100.0%; Pred. No.
 1.1%; Score 15; DB 1; Lo
100.0%; Pred. No. 1.2e+02;
 0;
0; Mismatches
 Score 14.8; DB 1; Length 18; Pred. No. 1e+02;
 0; Mismatches
 Mismatches
 DB 1; Le
. 1.2e+02;
 Length 20
 0; Indels
 Length 20;
 Indels
 Indels
0;
 0
 0;
Gaps
 Gaps
 Gaps
 0
 0
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Sequence 2967, Application US/09969373

Sequence 2967, Application US/09969373

Patent No. US20020133852A1

GENERAL INFORMATION:
APPLICANT: Effertz, Roger J.
APPLICANT: Hauge, Brian M.
TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
PILE REFERENCE: 38-10(52679)A

CURRENT APPLICATION NUMBER: US/09/969,373

CURRENT FILING DATE: 2001-10-02

PRIOR APPLICATION NUMBER: US 09/754,853

PRIOR APPLICATION NUMBER: US 09/760,427

PRIOR PILING DATE: 2001-01-13

PRIOR PRILING DATE: 2001-05-15

NUMBER OF SEG ID NOS: 4593

SEG ID NO 2967
 ; SEQ ID NO 4219
; LENGTH: 19
; TYPE: DNA
; ORGANISM: Glycine max
US-09-969-373-4219
 APPLICANT: Effertz, Roger J.

APPLICANT: Hauge, Brian M.

FITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
FILE REFERENCE: 38-10(52679)A

CURRENT APPLICATION NUMBER: US/09/969,373

CURRENT FILING DATE: 2001-10-02

PRIOR APPLICATION NUMBER: US 09/754,853

PRIOR APPLICATION NUMBER: US 09/760,427

PRIOR APPLICATION NUMBER: US 09/760,427
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 S
 ; TYPE: DNA; ORGANISM: Glycine max US-09-969-373-2967
 US-09-969-373-2967/c
 S
 US-09-969-373-4219
 Sequence 4219, Application US/09969373 Patent No. US20020133852A1 GENERAL INFORMATION:
 Best Local Similarity
Matches 16; Conserv
 Query Match
 Matches
 Query Match
Best Local
 NUMBER OF SEQ ID NOS: 4593
 LENGTH: 19
 1.1%;
Local Similarity 88.9%;
les 16; Conservarium
 2192 TGAAAATAGCAGACTTTG 2209
 1879 GAGATGATGAAGATGATT 1896
 2546 ATCGAATTCTCACTCTCA 2563
 18 TGAAACCAGCAGACTTTG 1
N
 ATGGAATTCTCACACTCA 1
 Conservative
 1.1%;
 0
 0; Mismatches
 Score 14.8; DB 1;
Pred. No. 1.1e+02;
 Score 14.8; DB 1;
Pred. No. 1.1e+02;
 Mismatches
 DB 1;
 Length 19;
 Length 19;
 Indels
 Indels
 0
 0;
 Gaps
 Gaps
 0
 0
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RESULT 51

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 US-10-357-488-32
 Query Match
Best Local S
 SOFTWARE: PatentIn version 3.1 SEQ ID NO 32
 Matches
 -09-866-108-8374/c
 GENERAL INFORMATION:
 Publication No.
 Sequence 8374, Application US/09866108
 PRIOR
PRIOR
 APPLICANT:
 FILE REFERENCE: 782-indian
CURRENT APPLICATION NUMBER: US/10/357,488
CURRENT FILING DATE: 2003-02-04
PRIOR APPLICATION NUMBER: 260/MAS/2002
PRIOR FILING DATE: 2002-04-08
NUMBER OF SEQ ID NOS: 37
 APPLICANT: Centre For DNA Fingerprinting and Diagnostics
TITLE OF INVENTION: No. US20030194730A1el FISSR-PCR primers and markers and a method
TITLE OF INVENTION: primers and markers for identifying genetic constitution and bre
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 PRIOR APPLICATION NUMBER: US 60/207,456
 FILE REFERENCE: AEOMICA-7
 APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICANT:
 PPLICANT: GU, Yizhong
 OTHER INFORMATION: A novel FISSR-PCR primer for genotyping eukaryotes
 ORGANISM: Artificial Sequence
 TYPE: DNA
 FEATURE:
 ENGTH: 16
 APPLICATION NUMBER: US 60/236,359 FILING DATE: 2000-09-27
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00662
 PILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 APPLICATION |
FILING DATE:
 APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00667
 APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04
 FILING DATE: 2000-05-26
 FILING DATE:
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00666
 APPLICATION NUMBER: PCT/US01/00661
 2473 ATGATGATGAGGGACT 2488
 Similarity
 CATION NUMBER: PCT/US01/00665
 ATGATGATGATGGACT 16
 CHEN, Wensheng
 HANZEL,
 PENN, Sharron
 Application US/10357488
 Conservative
 Yonggang
 2001-01-30
 2001-01-30
 2001-01-30
 David
 2001-05-25
 1.0%; Score 14.4;
93.8%; Pred. No. 93
 . ი
PCT/US01/00670
 Mismatches
 93;
 DB 1;
 IN HUMAN HEART AND MUSCLE
 Length 16;
 Indels
 0;
 Gaps
 0
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; SOPTWARE: Aeomica Sequence Listing Engine
; SEQ ID NO 8374
; LENGTH: 17
; TYPE: DNA
; ORGANISM: Homo Bapiens
US-09-866-108-8374
 RESULT 53
US-09-866-108-8375/c
 밁
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NUMBER OF SEQ ID NOS:
SOFTWARE: Aeomica Sequ
SEQ ID NO 8375
 Sequence 8375, Application US/09866108
Patent No. US20020048800A1
GENERAL INFORMATION:
 Matches
 Query Match
 APPLICANT:
 PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
 PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/0066
 PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: PCT PRIOR FILING DATE: 2001-01-30
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
 PRIOR APPLICATION NUMBER: US 60/234,687
 PRIOR
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 APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
 PRIOR FILING DATE: 2000-09-21
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
 RIOR FILING DATE:
 RIOR
 RIOR FILING DATE: 2001-01-30
 1.0%;
Local Similarity 93.8%;
es 15; Conservation
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00665 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00667
 FILING DATE: 2001-01-30
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER:
 FILING DATE:
 1506 CCAGCCGGCTGTGCAC 1521
 17
 CATION NUMBER: PCT/US01/00663
 PENN, Sharron G. HANZEL, David K.
 CHEN, Wensheng
 RANK, David R.
 Yizhong
 Yonggang
 2001-01-30
 2001-01-30
 2001-01-30
 PCT/US01/00670
 PCT/US01/00666
 PCT/US01/00664
 Listing Engine
 Score 14.4;
Pred. No. 1e
 Mismatches
 DB 1;
 Ä
 HUMAN HEART
 Length 17;
 Indels
 AND MUSCLE
 0
 Gaps
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 ; ORGANISM: Homo sapiens
US-09-866-108-8375
 US-09-866-108-9001
 RESULT 54
 Query Match
Best Local Similarity
Matches 15; Conserv
 TYPE: DNA.
ORGANISM: Homo sapiens
-09-866-108-9001
 SEQ ID NO 9001
 Matches
 Query Match
Best Local Similarity
 APPLICANT:
 Patent No.
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence
 PRIOR FILING DATE:
 FILE REFERENCE: AEOMICA-7
 PITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga
 APPLICATION NUMBER: US 60/266,860 FILING DATE: 2001-02-05
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
 FILING DATE: 2000-10-04
 APPLICATION NUMBER:
 APPLICATION NUMBER: US 60/236,359
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00670
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00665
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00661
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00668
 FILING DATE:
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00669
 1506 CCAGCCGGCTGTGCAC 1521
 1505 GCCAGCCGGCTGTGCA 1520
 ILING DATE: 2001-01-30
 LING DATE:
 16 CCAGCTGGCTGTGCAC
 001, Application US/09866108
US20020048800A1
GCCAGCCGGCGGTGCA 17
 RANK, David R. CHEN, Wensheng
 PENN, Sharron G. HANZEL, David K.
 Conservative
 Conservative
 2000-09-21
 2001-01-30
 2001-01-30
 2001-01-30
 2000-09-27
 2001-01-30
 2001-01-30
 1.0%;
 1.0%;
 PCT/US01/00666
 Listing
 Score 14.4; D
Pred. No. 1e+0
0; Mismatches
 ٥.
 Score 14.4; DB
Pred. No. 1e+02;
0; Mismatches
 Engine
 1e+02;
 BB
 ۳.
 IN HUMAN HEART AND MUSCLE
 1;
 Length 17;
 Length 17;
 0
 0,
 Gaps
 Gaps
 0
 0
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밁
 S
RESULT 56
US-09-848-754A-2482
 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-9002
 PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
 RESULT 55
US-09-866-108-9002
 Sequence 9002, Application US/09866108 Patent No. US20020048800A1
 Best Local Similarity
Matches 15; Conserv
 Query Match
 SEQ ID NO 9002
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 APPLICANT:
 APPLICANT: GU,
 SOFTWARE: Aeomica Sequence Listing Engine
 PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
 ENGTH:
 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
 FILING DATE: 2000-09-21
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
 APPLICATION NUMBER: US
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663
 1505 GCCAGCCGGCTGTGCA 1520
 INFORMATION:
 ь.
 GCCAGCCGGCGGTGCA 16
 JI, Yonggang
PENN, Sharron
HANZEL, David
 CHEN, Wensheng
 Conservative
 Yizhong
 David R
 2001-05-25
 1.0%;
 <u>بر</u>
 60/266,860
 0;
 Score 14.4; I
Pred. No. 1e+0
0; Mismatches
 4; DB 1;
1e+02;
 Length 17;
 Indels
 0
 Gaps
 0
```

Sequence 2482, Application US/09848754A Publication No. US20030073207A1 GENERAL INFORMATION: APPLICANT: Ribozyme Pharmaceuticals, II TITLE OF INVENTION: Levels of Epiderma

Pharmaceuticals, Inc

Enzymatic Nucleic Acid Treatment of Diseases Levels of Epidermal Growth Factor Receptors

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 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-238-700-3390
 RESULT 57
US-10-238-700-3390/c
 밁
 8
 ; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-848-754A-2482
 US-10-084-839-3116/c
 FILE REFERENCE: MBHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
SOFTWARE: PatentIn version 3.0
SEQ ID NO 2482
 GENERAL INFORMATION:
 Query Match 1.0%;
Best Local Similarity 93.8%;
Matches 15; Conservative
 SEQ ID NO 3390
 GENERAL INFORMATION:
 Sequence 3390, Application US/10238700 Publication No. US20030153521A1
 Sequence 3116, Application US/10084839 Publication No. US20030186238A1
 Matches
 Query Match 1.0%;
Best Local Similarity 68.8%;
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
 PRIOR APPLICATION NUMBER: US 60/318,471
PRIOR FILING DATE: 2001-09-10
NUMBER OF SEQ ID NOS: 4666
SOFTWARE: Patentin version 3.0
 APPLICANT:
 PRIOR APPLICATION NUMBER: PCT/US 02/16840 PRIOR FILING DATE: 2002-05-29
 APPLICANT:
 APPLICANT:
 ENGTH: 17
 1722 GGGCAAGCCCCTGGGA 1737
 2317 CATCAGAGTGATGTCT 2332
 11; Conservative
 16 GGGCAAGCCCCTTGGA 1
 N
 CÁCCAGAGUGAUGUCU 17
Lymaicheva, Natalie E. Ma, WuPo
 Allawi, Hatim
 Kwiatkowski, Jr., Ro
Lukowiak, Andrew A.
 Eis, Peggy S.
Hall, Jeff G.
 Chehak, LuAnne
Curtis, Michelle L.
 Bartholomay, Christian T.
 Argue,
 Lyamichev, Victor
 Kaiser, Michae
 Ji, Lin
 Wave Technologies
 Brad T.
 0;
 Robert
 Score 14.4; DB 1; Length 17; Pred. No. 1e+02;
 Score 14.4; DB Pred. No. 1e+02;
 Mismatches
 Mismatches
 DB 1;
 Length 17;
 Indels
 0
 0;
 Gaps
 Gaps
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 US-10-230-006-790
 RESULT 60
 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-230-006-789
 US-10-230-006-789
 US-10-084-839-3116
 Sequence 790, Application US/10230006 Publication No. US20030191077A1 GENERAL INFORMATION:
 SEQ ID NO 789
LENGTH: 17
 CURRENT APPLICATION NUMBER: US/10/084,839
CURRENT FILING DATE: 2002-02-26
NUMBER OF SEQ ID NOS: 4004
SOFTWARE: PatentIn version 3.1
SEQ ID NO 3116
LENGTH: 17
TYPE: DNA
 Matches
 Query Match
 GENERAL INFORMATION:
 Sequence 789, A Publication No.
 Matches
 Query Match 1.0%;
Best Local Similarity 93.8%;
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Fosnaugh, Kathy
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: METHOD AND REAGENT FOR THE TREATMENT OF ASTHMA AND ALLERGIC CONDITILE REFERENCE: 400/056 (MBHB01-1110)
 FILE REFERENCE: 400/056 (MBHB01-1110)
CURRENT APPLICATION NUMBER: US/10/230,006
CURRENT FILING DATE: 2002-11-18
 APPLICANT:
 PRIOR APPLICATION NUMBER: US 60/315,315 PRIOR FILING DATE: 2001-08-28
 APPLICANT: FOSNAUGH, KATHY
APPLICANT: MCSWiggen, Jim
TITLE OF INVENTION: METHOD AND REAGENT FOR THE TREATMENT OF ASTHMA AND ALLERGIC CONDI
 APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Fognaugh, Kathy
 SOFTWARE: PatentIn version 3.0
 NUMBER OF SEQ ID NOS: 2678
 FILE REFERENCE: FORS-06666
 APPLICANT: Vedvik, Kevin L.
TITLE OF INVENTION: RNA Detection Assays
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICANT:
 OTHER INFORMATION: Synthetic
 FEATURE:
 ORGANISM: Artificial Sequence
 Local
 1992 AGAATACCTCCGAGCC 2007
 1572 GTCCAGCTCCTCCATG 1587
 17 GTCCAGCTCCTCCCTG 2
 15;
 14;
 2 AGAAGACCUCCGAGCC 17
 Similarity
 Skrzypczynski, Zbigniew
Takova, Tsetska Y.
 Olson, Sarah M.
Olson-Munoz, Marilyn C.
 Thompson, Lisa C.
 Schaefer, James J.
 Application US/10230006
 Conservative
 Conservative
 US20030191077A1
 1.0%;
 <u>.</u>
 Score 14.4; DB 1; Length 17; Pred. No. 1e+02;
 Score 14.4; DB Pred. No. 1e+02;
 Mismatches
 Mismatches
 ۲,
 Indels
 0
 0
 0
 0
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANTION: Epidermal Growth Factor Receptors
FILE REFERENCE: MBH800-958-K (400/064)
CURRENT APPLICATION NUMBER: US/10/277,494
CURRENT FILING DATE: 2002-10-21
NUMBER OF SEQ ID NOS: 446
SOFTWARE: Patentin version 3.0
SEQ ID NO 46
LENGTH: 15
TYPE: RNA
ORGANISM: Homo sapiens
US-10-277-494-46
 RESULT 62
US-10-277-494-46
 В
 δ
 멍
 á
 ; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-230-006-790
 US-10-277-494-41
 RESULT 61
 US-10-277-494-41
 Sequence 41, Application US/10277494
Publication No. US20030186909A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals,
APPLICANT: McSwiggen, Jim
 Sequence 46, Application US/10277494 Publication No. US20030186909A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 41 LENGTH: 15
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 790
Query Match 1.0%;
Best Local Similarity 71.4%;
Matches, 10; Conservative
 Query Match 1.0%;
Best Local Similarity 71.4%;
Matches 10; Conservative
 Matches
 Query Match
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or conditions Related To Level
TITLE OF INVENTION: Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-K (400/064)
CURRENT APPLICATION NUMBER: US/10/277,494
 CURRENT FILING DATE: 2002-10-21
NUMBER OF SEQ ID NOS: 446
 PRIOR APPLICATION NUMBER: US 60/315,315
PRIOR FILING DATE: 2001-08-28
NUMBER OF SEQ ID NOS: 2678
 CURRENT FILING DATE:
 TYPE: RNA
ORGANISM: Homo sapiens
 Local
 2320 САСАСТСАТСТСТС 2333
 1992 AGAATACCTCCGAGCC 2007
 _
 14;
 N
 n 1.0%;
Similarity 87.5%;
14; Conservative
 AGAAGACCUCCGAGCC 16
 CAGAGUGAUGUCUG 15
 2002-11-18
 ; Score 14; DB; Pred. No. 93; 4; Mismatches
 4.
 Score 14.4; DB 1; Length 17; Pred. No. 1e+02; 1; Mismatches 1; Indels
 Score 14;
Pred. No.
 Mismatches
 Inc.
 DB
93;
 DB 1;
 1;
 0
 0
 Length 15;
 Length 15;
 Indels
 Indels
 0,
 0,
 0;
 Gaps
 Gaps
 Gaps
 ٥,
 0
 0
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SOFTWARE: Acomica Sequenc
SEQ ID NO 6888
LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
 8
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 S
 US-09-866-108-6888
 US-09-866-108-6888/c
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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 PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR APPLICATION NUMBER: DCT-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga
APPLICANT: PENN, Shar
APPLICANT: HANZEL, Da
 NUMBER OF SEQ ID NOS: 15752
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR FILING DATE: 2001-01-30
 FILE REFERENCE: AEOMICA-7
 PRIOR FILING DATE: 2001-02-05
 APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 PRIOR APPLICATION NUMBER: US 60/266,860
 APPLICANT:
 PRIOR APPLICATION NUMBER: PCT/US01/00663
 Local Similarity
 1358 CGCCTGGAAGAGAA 1371
 2321 AGAGTGATGTCTGG 2334
 INFORMATION:
17
 14;
 US20020048800A1
 AGAGUGAUGUCUGG
 PENN, Sharron
HANZEL, David
 RANK, David R. CHEN, Wensheng
 Conservative
 Application US/09866108
 Yonggang
 Sequence Listing Engine
 1.0%; Score 14;
100.0%; Pred. No.
 , ი
 14
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 Mismatches
 DB 1;
 1.2e+02;
 .,
 Length 17;
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 Gaps
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RESULT 64 US-09-866-108-6889/c

GENERAL INFORMATION:

Sequence 6889, Application US/09866108 Patent No. US20020048800A1

CURRENT FILING DATE:

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 ; ORGANISM: Homo sapiens
US-09-866-108-6889
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 Sequence 6890, Application US/09866108 Patent No. US20020048800A1
 APPLICANT:
APPLICANT:
APPLICANT:
 Matches
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Best Local Similarity
 -09-866-108-6890/c
 APPLICANT:
 APPLICANT: GU, Yizhong
 SOFTWARE: Aeomica Sequence Listing Engine EQ ID NO 6889
 NUMBER OF SEQ ID NOS: 15752
 CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
 CURRENT APPLICATION NUMBER: US/09/866,108
 ENGTH:
 PLICANT: SHANNON, MAIK
TLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
 1358 CGCCTGGAAGAGAA 1371
 APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00661
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: US 60/236,359
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00667
 APPLICATION NUMBER: PCT/US01/00666
 FILING DATE: 2000-09-27
 INFORMATION:
APPLICATION NUMBER: US/09/866,108
 16 CGCCTGGAAGAGAA 3
 T: JI, Yonggang
T: PENN, Sharron G.
T: HANZEL, David K.
T: RANK, David R.
T: CHEN, Wensheng
TI: SHANNON, Mark
INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 LING DATE: 2001-01-30
 LING DATE: 2001-01-30
 LING DATE: 2001-01-30
 LING DATE: 2001-01-30
 14;
 LING DATE:
 LING DATE: 2000-05-
 NG DATE: 2001-01-30
CATION NUMBER: PCT/US01/00663
 CHEN, Wensheng
 RANK, David R
 Conservative 0;
 AEOMICA-
 Yonggang
 2001-01-30
 2001-01-30
 2001-01-30
 1.0%;
 Score 14;
 Mismatches
 1.2e+02;
nes 0;
 DB 1;
 Length 17;
 0,
 Gaps
 0
```

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 Ś
 ; ORGANISM: Homo sapiens
US-09-866-108-6890
 US-09-866-108-6891/c
 RESULT 66
 Best Loc
Matches
 GENERAL
 Sequence 6891, Application US/09866108
 Query Match
 Patent No. US20020048800A1
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
 APPLICANT:
 APPLICANT: GU, Yizhong APPLICANT: JI, Yongga
 PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
 APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 PRIOR APPLICATION NUMBER: US 60/236,359 PRIOR FILING DATE: 2000-09-27
 FILE REFERENCE: AEOMICA-7
 CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 NUMBER OF SEQ ID NOS: 15752
 PRIOR FILING DATE: 2000-09-27
 PRIOR APPLICATION NUMBER: GB 24263.6 PRIOR FILING DATE: 2000-10-04
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: US 60/236,359
 LENGTH:
 Local
 APPLICATION NUMBER:
 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00666 FILING DATE: 2001-01-30
 1358 CGCCTGGAAGAGAA 1371
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00662
 FILING DATE: 2001-01-30
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER:
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00667
 INFORMATI
 15
 LING DATE:
 LING DATE:
 Similarity
 Aeomica Sequence Listing Engine
 NG DATE: 2001-01-30
ICATION NUMBER: PCT/US01/00661
 CATION NUMBER: PCT/US01/00663
 JI, Yonggang
PENN, Sharron G.
HANZEL, David K.
 DATE:
 Conservative (
 2001-01-30
 2001-01-30
 001-01-30
 -01-30
PCT/US01/00667
 PCT/US01/00666
 PCT/US01/00664
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 Score 14;
; Pred. No.
 DB 1;
 1.2e+02;
 Length 17;
 ..
 Gaps
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RESULT 68
US-09-776-474-1034
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 8
 ; SEQ ID NO 2483
; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-848-754A-2483
 밁
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 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-6891
 US-09-848-754A-2483
 RESULT 67
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MHHB00-958-I (400/018)
CURRENT APPLICATION UNMEBER: US/00/848,754A
CURRENT FILING DATE: 2001-05-03
 Sequence 1034, Application US/09776474 Publication No. US20030087847A1 GENERAL INFORMATION:
 Query Match
Best Local :
 Sequence 2483, Application US/09848754A Publication No. US20030073207A1
 Query Match 1.0%; Score 14; DB 1; L
Best Local Similarity 100.0%; Pred. No. 1.2e+02;
Matches 14; Conservative 0; Mismatches 0;
 Matches
 PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
 PRIOR
PRIOR
 PRIOR
PRIOR
APPLICANT: Ribozyme APPLICANT: Jarvis,
 NUMBER OF SEQ ID NOS: 9645
SOFTWARE: PatentIn version 3.0
 SOFTWARE: Aeomica Sequence Listing Engine
 ID NO 6891
ENGTH: 17
 ENGTH:
 Local Similarity 71.4%;
nes 10; Conservative
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00670
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
 2321 AGAGTGATGTCTGG 2334
 1358 CGCCTGGAAGAGAA 1371
 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
 FILING DATE:
 14 CGCCTGGAAGAGAA 1
 AGAGUGAUGUCUGG 14
 2001-01-30
 Pharmaceuticals, Thale
 Score 14; DB
Pred. No. 1.20
4; Mismatches
 .2e+02;
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 DB 1; Length 17;
 Length 17;
 Indels
 0;
 0
 Gaps
 Gaps
 0;
 0
 US-09-776-474-1036
 RESULT 70
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 US-09-776-474-1035
 US-09-776-474-1035
 OTHER INFORMATION: Description of Artificial Sequence: US-09-776-474-1034
 Query Match
Best Local Similarity
Matches 12; Conserv
 SOFTWARE: PatentIn version 3.0
SEQ ID NO 1034
LENGTH: 17
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 1035
 GENERAL INFORMATION
 Publication No.
 Query Match
 Matches
 APPLICANT:
APPLICANT:
APPLICANT:
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Inhibition of Checkpoint Kinase-1
TITLE OF INVENTION: Enzyme
FILE REFERENCE: MeHB00-955-A (400/008)
CURRENT APPLICATION NUMBER: US/09/776,474
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 2992
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Jarvis, Thale APPLICANT: Booher, Robert APPLICANT: Holman, Patricia APPLICANT: Fattaey, Ali
 PRIOR FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 2992
 TITLE OF INVENTION: Method and Reagent fo:
TITLE OF INVENTION: METHOD and Reagent fo:
FILE REFERENCE: MBHB00-955-A (400/008)
CURRENT APPLICATION NUMBER: US/09/776,474
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,983
 APPLICANT:
 FEATURE: OTHER INFORMATION: Description of Artificial Sequence:
 ORGANISM: Artificial Sequence
 LENGTH: 17
TYPE: RNA
 FEATURE:
 TYPE: RNA
 ORGANISM: Artificial Sequence
 Local Similarity les 12; Conserv
 1730 CCCTGGGAGAAGGT 1743
 1730 CCCTGGGAGAAGGT 1743
w
 4
 CCCUGGGAGAAGGU 17
 CCCUGGGAGAAGGU 16
 McSwiggen, Jim
 Holman, Patricia
 Booher, Robert
 Fattaey, Ali
 Application US/09776474 US20030087847A1
 Conservative
 Conservative
 1.0%;
 1.0%;
85.7%;
 2;
 2
 Score 14; DB 1; Lo
Pred. No. 1.2e+02;
2; Mismatches 0;
 Score 14; I
Pred. No. 1
 Mismatches
 for the Inhibition of Checkpoint Kinase-1
 DB 1;
1.2e+02;
 Length 17
 Length 17;
 Nucleic Acid
 Nucleic
 0
 0
 Gaps
 Gaps
 0
 0;
 (CHK
 (CHK
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Sequence 1036, Application US/09776474 Publication No. US20030087847A1 GENERAL INFORMATION:

APPLICANT: Ribozyme

Pharmaceuticals, Inc

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S
 ; OTHER INFORMATION: Description of Artificial Sequence:
US-09-776-474-1036
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Inhibition of Checkpoint Kinase-1 (CHR
TITLE OF INVENTION: Enzyme
FILE REFERENCE: MBHB00-955-A (400/008)
CURRENT FILING NUMBER: US/09/776,474
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 2992
SOFTWARE: Patentin version 3.0
SEQ ID NO 1036
 Matches
 -09-866-108-6312
 Query Match
 APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
 CURRENT APPLICATION NUMBER: US/09/866,108
 ORGANISM: Artificial Sequence FEATURE:
 TYPE: RNA.
 ENGTH: 17
 OR APPLICATION NUMBER: PCT/US01/00666
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00667
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00664
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00669
 Local Similarity
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
 APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04
 APPLICATION NUMBER: US 60/207,456 FILING DATE: 2000-05-26
 APPLICATION NUMBER: US 60/236,359 FILING DATE: 2000-09-27
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 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00661
 1730 CCCTGGGAGAAGGT 1743
 INFORMATION
 6312, Application US/09866108
5. US20020048800A1
 12; Conservative
 1 CCCUGGGAGAAGGU 14
ICATION NUMBER: PCT/US01/00670
 PENN, Sharron G.
HANZEL, David K.
RANK, David R.
 CHEN, Wensheng
 Fattaey, Ali
 Yonggang
 izhong
 2001-01-30
 Patricia
 Robert
 2001-05-25
 1.0%;
 Score 14; DB 1;
Pred. No. 1.2e+02;
 Mismatches
 IN HUMAN HEART AND MUSCLE
 0
 Length 17
 Indels
 Nucleic Acid
 0
 Gaps
 0
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 US-09-866-108-7673/c
 ; ORGANISM: Homo sapiens US-09-866-108-6312
 PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aecmica Sequence Listing Engine
SEQ ID NO 7673
 Best Local Similarity
Matches 15; Conserv
 SEQ ID NO 6312
 Patent No.
 Query Match
 APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: ABOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
CURRENT FILING DATE: 2001-05-25
 PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
NUMBER OF SEQ ID NOS: 15752
 LENGTH: 17
TYPE: DNA
 SOFTWARE: Aeomica Sequence Listing Engine
 PRIOR APPLICATION NUMBER: US 60/266,860
 PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 PRIOR APPLICATION NUMBER: US 60/234,687
 PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00667 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
 PRIOR APPLICATION NUMBER: GB 24263.6
ORGANISM: Homo sapiens
 APPLICATION NUMBER: US 60/234,687
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00662
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00668
 2090 GCACCTACCAGCTGGCC 2106
 7673, Application US/09866108
5. US20020048800A1
 CHEN, Wensheng
 PENN, Sharron G.
HANZEL, David K.
 JI, Yonggang
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 2001-01-30
 2000-09-21
 2001-01-30
 2001-01-30
 2001-01-30
 2000-09-27
 2001-01-30
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88.2%;
 0; Mismatches -
 Score 13.8;
Pred. No. 1
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 DB 1;
 Length 17;
 Indels
 0
 Gaps
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 US-09-866-108-9075
 8
 ; TYPE: DNA; Homo sapiens; ORGANISM: Homo sapiens US-09-866-108-9075
 US-09-866-108-7673
 SEQ ID NO 9075
LENGTH: 17
 Matches
 Query Match
 Matches
 Query Match
Best Local
 Sequence 9075, Application US/09866108 Patent No. US20020048800A1
 Best Local Similarity
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 SOFTWARE: Aeomica Sequence Listing Engine
 NUMBER OF SEQ ID NOS: 15752
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: US 60/207,456
 FILE REFERENCE: AEOMICA-7
 PPLICANT: SHANNON, Mark
ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04
 APPLICATION NUMBER: US 60/266,860 FILING DATE: 2001-02-05
 APPLICATION NUMBER: US 60/236,359
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00667
 APPLICATION NUMBER: PCT/US01/00666 FILING DATE: 2001-01-30
 FILING DATE: 2000-09-27
 1397 ACCTGGAGATAGCCATT 1413
 FILING DATE: 2001-01-30
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00663 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00665
 2658 TTCTGTTTTTTCTCCAG 2674
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00668
 INFORMATION
 ILING PATE: 2001-01-30
 [LING DATE:
 LING DATE:
 17 TTCTGCTTCTTCTCCAG
 LING DATE:
 LING DATE:
 15;
 Similarity
ACCTGGAGACATCCATT 17
 JI, Yonggang
PENN, Sharron G.
HANZEL, David K.
 RANK, David R. CHEN, Wensheng
 Conservative
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 2001-01-30
 2000-09-21
 2001-01-30
 2001-01-30
 2001-01-30
 2000-05-26
 2001-01-30
 88.2%;
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 Score 13.8; DB 1;
Pred. No. 1.2e+02;
0; Mismatches 2;
 Score 13.8; DB 1;
Pred. No. 1.2e+02;
 Mismatches
 2;
 Length 17;
 Length 17;
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 Gaps
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 0
 0
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 US-09-853-526-84/c
 ; NAME/KEY: misc_feature
; LOCATION: (1)...(17)
; OTHER INFORMATION: sequencing oligonucleotide moPGrace5R444
US-09-901-484A-84
 RESULT 74
US-09-901-484A-84/c
 PRIOR APPLICATION NUMBER: US 08/996,306
PRIOR FILING DATE: 1997-12-22
PRIOR APPLICATION NUMBER: US 60/099,658
PRIOR FILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: US 09/218,207
PRIOR APPLICATION NUMBER: US 09/338,907
PRIOR APPLICATION NUMBER: US 09/338,907
PRIOR FILING DATE: 1999-06-23
PRIOR FILING DATE: 1999-06-23
 Patent No. Ubzv...
 SOFTWARE: P
 Sequence 84, Application US/09901484A Patent No. US20020119460A1
 SEQ ID NO 84
 Sequence
 Matches
 Query Match
Best Local Similarity
 Patent No.
 FILE REFERENCE: GENSET.18CP1CP
CURRENT APPLICATION NUMBER: US/09/853,526
CURRENT FILING DATE: 2001-05-11
PRIOR APPLICATION NUMBER: 09/338,907
PRIOR FILING DATE: 1999-06-23
PRIOR APPLICATION NUMBER: 08/996,306
 APPLICANT: Ilya, Chumano, APPLICANT: Bougueleret, Lydie TITLE OF INVENTION: PROSTATE CFTITLE OF ENERGINES CHESET, IBCENCE; GENSET, IBCENCE, US
 NUMBER OF SEQ ID NOS: 578
 PRIOR APPLICATION NUMBER: US 09/853,526
PRIOR FILING DATE: 2001-05-11
 FILE REFERENCE: GEN-T111XC3D2
CURRENT APPLICATION NUMBER: US/09/901,484A
CURRENT FILING DATE: 2001-07-09
 NUMBER OF SEQ ID NOS: 578
 PRIOR APPLICATION NUMBER: 09/218,207
PRIOR FILING DATE: 1998-12-22
 PRIOR FILING DATE: 1997-12-22
PRIOR APPLICATION NUMBER: 60/099,658
 APPLICANT: Cohen, Daniel
APPLICANT: Blumenfeld, Marta
 SOFTWARE:
 PRIOR FILING DATE: 1998-09-09
 APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: Prostate Cancer Gene
 TYPE: DNA
 ORGANISM: Mus musculus FEATURE:
 TYPE: DNA
FEATURE:
 ORGANISM: Mus Musculus
 ENGTH:
 LENGTH:
 2092 ACCTACCAGCTGGCCAG
 INFORMATION:
 84, Application US/09853526
o. US20020165345A1
 17 ACCTACCTGCTGGCCTG 1
 15;
 PatentIn version 3.1
 Patent.pm
 Blumenfeld,
 Chumakov, Ilya
 Conservative
 Daniel
 1.0%;
 Marta
 2108
 CANCER GENE
 °;
 Score 13.8; DB 1
Pred. No. 1.2e+02
 Mismatches
 Length 17
 Indels
 0
 Gaps
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 US-09-825-805-384/c
 RESULT 77
 OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid
US-09-864-785-2863
 ; OTHER INFORMATION: sequencing oligonucleotide moPGrace5R444
US-09-853-526-84
 US-09-864-785-2863
 RESULT 76
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of NF-Kappa B
FILE REFERENCE: 400/022 (MBHB00-812-D)
CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 3929
SOFTWARE: Patentin version 3.0
SEQ ID NO 2863
 Sequence 384, Application US/09825805 Publication No. US20030004122A1
 Sequence 2863, Application US/09864785 Patent No. US20020177568A1
 APPLICANT:
 GENERAL INFORMATION
 Matches
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 Matches
 Query Match
 PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 09/476,387
PRIOR FILING DATE: 1999-12-30
PRIOR APPLICATION NUMBER: 09/474,432
PRIOR FILING DATE: 1999-12-29
PRIOR FILING DATE: 1999-12-29
 APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleotide Triphosphate and their Incorporation into Oligonucleot
PILE REFERENCE: MBHB00-831-F (400/009)
CURRENT APPLICATION NUMBER: US/09/825,805
CURRENT FILLING DATE: 2001-09-27
CURRENT FILLING DATE: 2001-09-27
ERIOR APPLICATION NUMBER: 09/578,223
 APPLICANT: Ribozyme Pharmaceuticals,
APPLICANT: Beigelman, Leo
APPLICANT: Beaudry, Amber
APPLICANT: Karpeisky, Alex
APPLICANT: Adamic, Jasenka Matulic
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Stinchcomb, Dan
APPLICANT: Draper, Ken
 TYPE: RNA
ORGANISM: Artificial Sequence
 NAME/KEY: misc_feature LOCATION: 1..17
 FEATURE:
 LENGTH: 17
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APPLICATION NUMBER: 09/301,511
FILING DATE: 1999-04-28
 2004 AGCCCGGAGGCCACCCG 2020
 2092 ACCTACCAGCTGGCCAG 2108
 15; Conservative
 17 ACCTACCTGCTGGCCTG 1
 AGCCCAGAGGCCCCCCG 17
 Sweedler, Dave
 1.0%;
ilarity 88.2%;
Conservative
 1.0%;
 <u>,</u>
 0
 Score 13.8;
Pred. No. 1.
 Score 13.8;
Pred. No. 1
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 Mismatches
 1.2e+02;
 2e+02
 B
 DB 1;
 1;
 Length 17;
 Length 17;
 Indels
 0
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 Gaps
 Gaps
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 ; TYPE: RNA; Homo sapiens US-09-825-805-777
 RESULT 78
US-09-825-805-777
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 US-09-825-805-384
 Query Match
Best Local Similarity
Matches 12; Conserv
 NUMBER OF SEQ ID NOS: 1558
SOFTWARE: PatentIn version 3.0
SEQ ID NO 777
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals,
 Sequence 777, A Publication No.
 Best Local Similarity 88.2%; Matches 15; Conservative
 SOFTWARE: Pat
SEQ ID NO 384
 Query Match
 PRIOR APPLICATION NUMBER: 60/083,727
PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: 60/064,866
PRIOR FILING DATE: 1997-11-05
NUMBER OF SEQ ID NOS: 1558
 CURRENT APPLICATION NUMBER: US/09/825,805
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: 09/578,223
PRIOR FILING DATE: 2000-05-23
 PRIOR APPLICATION NUMBER: 09/476,387
PRIOR FILING DATE: 1999-12-30
PRIOR APPLICATION NUMBER: 09/474,432
 APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleotide Triphosphate and their Incorporation into Oligonucleot
FILE REFERENCE: MBHB00-831-F (400/009)
 APPLICANT:
 APPLICANT:
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: 60/064,866
 PRIOR FILING DATE: 1998-04-29
 PRIOR APPLICATION NUMBER: 60/083,727
 PRIOR FILING DATE: 1998-11-04
 PRIOR APPLICATION NUMBER: 09/186,675
 PRIOR APPLICATION NUMBER: 09/301,511 PRIOR FILING DATE: 1999-04-28
 PRIOR FILING DATE: 1999-12-29
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 LENGTH: 17
 2269 CCAGTCAAGTGGATGGC 2285
 1920 TCTTCTTGGAGCCTGCA 1936
 17
1 CCCAUCAAGUGGAUGGC
 PatentIn version 3.0
 Beigelman, Leo
Beaudry, Amber
 Karpeisky, Alex
Adamic, Jasenka Matulic
 Sweedler,
 Application US/09825805

DS20030004122A1
 Conservative
 1997-11-05
 Dave
 1.0%;
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 0; Mismatches
 Score 13.8;
Pred. No. 1
 Score 13.8;
Pred. No. 1.
 Mismatches
 1.2e+02
 1.2e+02;
 DB 1;
 Length 17;
 Indels
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 Gaps
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RESULT 79 US-09-780-533A-844/c

Sequence 844, Application US/09780533A Publication No. US20030060611A1

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 RESULT 80
US-09-877-478-1296
 밁
 S
 US-09-780-533A-844
 Query Match
Best Local Similarity
Matches 15; Conserv
 SEQ ID NO 1296
 GENERAL INFORMATION:
 SOFTWARE: Pai
SEQ ID NO 844
 Matches
 Query Match
Best Local
 Sequence 1296, Application US/09877478 Publication No. US20030068301A1
 TYPE: RNA
ORGANISM: Hepatitis B virus
-09-877-478-1296
 APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MSHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
 NUMBER OF SEQ ID NOS: 6586 SOFTWARE: PatentIn version
 TITLE OF INVENTION: Method and Reagent for Inhibiting Hepatitis B Virus Replication FILE REFERENCE: MBHB0-845-H (400/029) CURRENT APPLICATION NUMBER: US/99/877,478 CURRENT FILING DATE: 2001-12-31
 NUMBER OF SEQ ID NOS: 6679
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: US 07/882,712 PRIOR FILING DATE: 1992-05-14
 APPLICANT:
 APPLICANT:
 APPLICANT: Ribozyme Pharmaceuticals, Inc
 APPLICANT: Ribozyme Pharmaceuticals, Inc
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 ENGTH: 17
 APPLICATION NUMBER: US 09/636,385
FILING DATE: 2000-08-09
APPLICATION NUMBER: US 09/696,347
 APPLICATION NUMBER: US 09/531,025 FILING DATE: 2000-03-20
 APPLICATION NUMBER: US 09/436,430
 FILING DATE: 2000-10-24
APPLICATION NUMBER: US 08/193,627
FILING DATE: 1994-02-07
APPLICATION NUMBER: US 08/433,993
 FILING DATE:
 APPLICATION NUMBER: US 08/434,504
 FILING DATE: 1995-05-04
 2191 ATGAAAATAGCAGACTT 2207
 1650 GCTGGCAGGGGTCTCCG 1666
 17 GCCGGCAGGGGTCCCCG 1
 Similarity
 PatentIn version 3.0
AUGAAAACAGGAGACUU 17
 McSwiggen, Jim
Morrissey, Dave
 McSwiggen, Jim
Chowrira, Bharat
 Blatt, Larry
 Draper,
 Conservative
 Conservative
 1999-11-08
 1995-05-04
 Larry
 Kenneth
 1.0%;
 70.6%;
 0;
 Score 13.8; DB 1;
Pred. No. 1.2e+02;
3; Mismatches 2;
 Score 13.8;
Pred. No. 1.
 Mismatches
 .2e+02
 В
 1;
 Length 17
 Length 17;
 Indels
 0
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 Gaps
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 RESULT 83
US-09-776-474-430
 US-09-848-754A-3492
 US-09-848-754A-3492
 US-09-848-754A-2480
 US-09-848-754A-2480
 RESULT 81
 Sequence 2480, Application US/09848754A Publication No. US20030073207A1 GENERAL INFORMATION:
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Jarvis, Thale
APPLICANT: Booher, Robert
APPLICANT: Holman, Patricia
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 3492
 SEQ ID NO 2480
 Sequence 430, A Publication No.
 Query Match
 Publication No. US20 GENERAL INFORMATION:
 Sequence 3492, Application US/09848754A
 Query Match
 Matches
 Matches
 TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment TITLE OF INVENTION: Levels of Epidermal Growth Facto FILE REFERENCE: MBHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
 TITLE OF INVENTION: Enzymatic Nucleic Acid Treat TITLE OF INVENTION: Levels of Epidermal Growth FILE REFERENCE: MBHB00-958-I (400/018) CURRENT APPLICATION NUMBER: US/09/848,754A CURRENT FILING DATE: 2001-05-03 NUMBER OF SEQ ID NOS: 9645
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent
TITLE OF INVENTION: Enzyme
 LENGTH: 17
TYPE: RNA
 APPLICANT:
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
 SOFTWARE: PatentIn version 3.0
 APPLICANT: Ribozyme
CURRENT APPLICATION NUMBER: US/09/776,474
 TYPE: RNA
ORGANISM: Homo sapiens
 ORGANISM: Homo sapiens
 LENGTH:
 Local Similarity
hes 12; Conserv
 Local Similarity
mes 12; Conserv
 2314 ACTCATCAGAGTGATGT 2330
 2269 CCAGTCAAGTGGATGGC 2285
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 17
 CCUAUCAAGUGGAUGGC
 ACCCACCAGAGUGAUGU
 Fattaey, Ali
 Application US/09776474
 Conservative
 Conservative
 US20030087847A1
 US20030073207A1
 ON: Enzyme
MBHB00-955-A (400/008)
 Pharmaceuticals, Inc.
Enzymatic Nucleic Acid Treatment of Diseases or Conditions
 1.0%;
 1.0%;
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 Score 13.8; DB 1;
Pred. No. 1.2e+02;
 Score 13.8;
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 Mismatches
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 Length 17;
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RESULT 85
US-09-780-164-921
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 Query Match
Best Local Similarity
Watches 12; Conserve
 TITLE OF INVENTION: McSwiggen, Jim
FILE REFERENCE: MBHB00,918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930,423
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARE: PATCHIN Version 3.0
SEQ ID NO 1394
LENGTH- 17
 밁
 S
 ; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo Sapiens
US-09-930-423-1394
 RESULT 84
US-09-930-423-1394
 ; OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid US-09-776-474-430
SEQ ID NO 921
LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 2992
SOFTWARE: PATENTIN VERSION 3.0
SEQ ID NO 430
 Sequence 921, Application US/09780164 Publication No. US20030092646A1
 Sequence 1394, Application US/09930423 Publication No. US20030092003A1
 GENERAL INFORMATION:
 Query Match
Best Local Similarity
 Matches
 PRIOR APPLICATION NUMBER: 2001-02-09
PRIOR FILING DATE: 2000-02-28
NUMBER OF TOTAL T
 FILE REFERENCE: 400/010
CURRENT APPLICATION NUMBER: US/09/780,164
CURRENT FILING DATE: 2001-02-09
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Inhibition of CD20
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
 NUMBER OF SEQ ID NOS: 2603
 SOFTWARE: PatentIn version 3.0
 CURRENT FILING DATE:
 TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
 ENGTH:
 2025 GGAGTACTCCTATGACA 2041
 2194 AAAATAGCAGACTTTGG 2210
 tch 1.0%;
al Similarity 64.7%;
ll; Conservative
 1 AAAAUCUCAGACUUUGG 17
 GGAGUACAACUAUGACA 17
 Conservative
 1.0%;
 2001-02-02
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 Score 13.8; DB 1;
Pred. No. 1.2e+02;
 Score 13.8;
Pred. No. 1
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 DB 1;
 DB 1;
 Length 17;
 Length 17;
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 Gaps
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밁
 US-09-745-237A-1394
 RESULT 87
US-09-745-237A-1394
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 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-780-164-1056
 RESULT 86
 멹
 US-09-780-164-1056
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 1394 LENGTH: 17
 Sequence 1394, Application US/09745237A Publication No. US20030143708A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0
SEQ ID NO 1056
LENGTH: 17
 Matches
 Best Local Similarity
Matches 12; Conserv
 Query Match
 Query Match
Best Local Similarity
 Query Match
 Sequence 1056, Application US/09780164
Publication No. US20030092646A1
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: 400/007 (MBHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
 FILE REFERENCE: 400/010
CURRENT APPLICATION NUMBER: US/09/780,164
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/185,516
PRIOR FILING DATE: 2000-02-28
 GENERAL INFORMATION:
 APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: Blatt, Larry
 LENGTH:
TYPE: RN
 NUMBER OF SEQ ID NOS: 2603
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Inhibition of CD20
 TYPE: RNA
ORGANISM: Homo sapiens
 Local Similarity
 2025 GGAGTACTCCTATGACA 2041
 1700 TTCCAAGAGATAAGCTG 1716
 1701 TCCAAGAGATAAGCTGA 1717
 13;
 1 UUCCAAGAGACAUGCUG
 1 UCCAAGAGACAUGCUGA 17
GGAGUACAACUAUGACA
 Conservative
 Conservative
 Conservative
 1.0%;
 1.0%;
 1.0%; Score 13.8; 170.6%; Pred. No. 1.20; ive 3; Mismatches
 17
 ; Score 13.8; D; Pred. No. 1.2e
 2
 Score 13.8; DB 1;
Pred. No. 1.2e+02;
 Mismatches
 8; DB 1;
1.2e+02;
 1.2e+02;
2;
 DB 1;
 DB 1;
 Length 17;
 Length 17;
 Length 17;
 Indels
 0,
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 Gaps
 Gaps
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US-10-061-201-122/c ; Sequence 122, Application US/10061201 ; Publication No. US20030166229A1

RESULT 88

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밁
 RESULT 89
 US-10-061-201-122
 Matches
 Query Match
 SEQ ID NO 122
 GENERAL INFORMATION:
 CURRENT APPLICATION NUMBER: US/10/061,20:
CURRENT FILING DATE: 2002-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
 CURRENT APPLICATION NUMBER: US/10/061,201
CURRENT FILING DATE: 2002-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00664
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: PCT/US01/00666
 ENERAL INFORMATION:
 SOFTWARE: Aeomica Sequence Listing Engine
 PRIOR APPLICATION NUMBER: PCT/US01/00667
 FILE REFERENCE:
 FILE REFERENCE:
 TYPE: DNA
ORGANISM: Homo sapiens
 NPPLICANT: Shannon,
 TLE OF INVENTION:
 ENGTH: 17
 Local
 APPLICATION NUMBER: US 60/328,205
 APPLICATION NUMBER: PCT/US01/00664 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
 FILING DATE:
 FILING DATE:
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00667
 APPLICATION NUMBER: US 09/864,761
 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00665 FILING DATE: 2001-01-30
 FILING DATE: 2001-05-
 APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663
 1564 TCGGCTGAGTCCAGCTC 1580
 APPLICATION NUMBER: US 09/864,761
 FILING DATE:
 1.0%;
1 Similarity 88.2%;
15; Conservation
 17 TCTGCTGAGTTCAGCTC 1
 LING DATE: 2001-01-30
 LING DATE:
 Application US/10061201
 US20030166229A1
NUMBER: US 60/328,205
 2001-10-10
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 2001-05-23
 2001-01-30
 HUMAN POSH-LIKE PROTEIN 1
 HUMAN POSH-LIKE PROTEIN
 2002-01-30
 0
 Score 13.8;
Pred. No. 1.
 Mismatches
 2e+02
 DB 1;
 Length 17;
 Indels
 0
 Gaps
 0
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밁
 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-339-782-153
 밁
 ; TYPE: DNA; Homo sapiens US-10-061-201-123
 US-10-339-782-153/c
 RESULT 90
 US-10-060-756A-145
 Query Match
Best Local Similarity
Matches 15; Conserv
 SEQ ID NO 153
LENGTH: 17
 GENERAL INFORMATION:
 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 123
 GENERAL INFORMATION
 Sequence 145, A Publication No.
 Matches
 Query Match
Best Local Similarity
 Sequence 153, A Publication No.
 CURRENT FILING DATE: 2003-01-1
NUMBER OF SEQ ID NOS: 495
SOFTWARE: PatentIn version 3.1
 FILE REFERENCE: 37-000110US
CURRENT APPLICATION NUMBER: US/10/339,782
 NUMBER OF SEQ ID NOS: 4162
 CURRENT APPLICATION NUMBER: US/10/060,756A
CURRENT FILING DATE: 2002-01-30
PRIOR APPLICATION UMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
 APPLICANT: Lynx Therapeutics, Inc. APPLICANT: Goodman, Laurie J
 PRIOR APPLICATION NUMBER: US 09/864,761 PRIOR FILING DATE: 2001-05-23
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
 PRIOR APPLICATION NUMBER: PCT/US01/00664 PRIOR FILING DATE: 2001-01-30
 APPLICANT: Zhang, Jian
TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
 TITLE OF INVENTION: Identification of Specific Biomarkers for Breast Cancer Cells
PRIOR APPLICATION NUMBER: US 60/327,898
PRIOR FILING DATE: 2001-10-09
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00668
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00669
 FILE REFERENCE: PB0177
 LENGTH:
 2532 GGTAGAAGACTTGGATC 2548
 1563 TTCGGCTGAGTCCAGCT 1579
 17 TTCTGCTGAGTTCAGCT 1
 17 GGTTGAAGACCTGGATC
 15;
 17
 Goodman, Laurie v
Bowen, Benjamin A
 Application US/10339782
o. US20030166026A1
 Application US/10060756A
 Conservative
 Conservative
 US20030046717A1
 1.0%;
 2003-01-08
 1.0%;
 <u>,</u>
 <u>,</u>
 Score 13.8;
Pred. No. 1.
 Score 13.8; DB 1
Pred. No. 1.2e+02
 Mismatches
 Mismatches
 1.2e+02;
 DB 1;
 Length 17;
 Length
 0,
 <u>,,</u>
 Gaps
 Gaps
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RESULT 93
US-10-163-552-303/c
 밁
 S
 ; TYPE: DNA; Homo sapiens; ORGANISM: Homo sapiens US-10-060-756A-1676
 US-10-060-756A-1676/c
 ફ
 US-10-060-756A-145
 Sequence 303, Application US/10163552 Publication No. US20030105051A1 GENERAL INFORMATION:
 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 145
 Query Match 1.0%;
Best Local Similarity 88.2%;
 SEQ ID NO 1676
 Matches
 Sequence 1676, Application US/10060756A
 Matches
 Best Local Similarity
 Query Match
FILE REFERENCE: MBHB01-1653-A (400/014)
CURRENT APPLICATION NUMBER: US/10/163,5
CURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 1997
 APPLICANT: Zhang, Jian
TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
FILE REFERENCE: PB0177
CURRENT APPLICATION NUMBER: US/10/060,756A
CURRENT FILING DATE: 2002-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
 ublication No.
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, Jim TITLE OF INVENTION: Nucleic acid treatment TITLE OF INVENTION: HER2
 SOFTWARE: Aeomica Sequence Listing Engine
 NUMBER OF SEQ ID NOS: 4804
 PRIOR APPLICATION NUMBER: PCT/US01/00664 PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2001-10-09
 PRIOR APPLICATION NUMBER: PCT/US01/00669
 LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
 ENGTH: 17
 APPLICATION NUMBER: PCT/USO1/00668 FILING DATE: 2001-01-30 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: US 60/327,898
 APPLICATION NUMBER: US 09/864,761 FILING DATE: 2001-05-23
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
 2186 ATGTGATGAAAATAGCA 2202
 1518 GCACAAGCTGACCAAAC 1534
 17 ATGTTATAAAATAGCA 1
 15;
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 US20030046717A1
 1.0%;
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 Score 13.8; D
Pred. No. 1.2e
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US-09-280-030-8
 US-09-280-030-8/c
 ; ORGANISM: Homo sapiens US-10-163-552-666
 RESULT 94
US-10-163-552-666
 ; TYPE: RNA; Homo sapiens; ORGANISM: Homo sapiens US-10-163-552-303
 TITLE OF INVENTION: DNAS ENCODING NEW FUSION PROTEINS AND PROCESSES FOR TITLE OF INVENTION: PREPARING USEFUL POLYPEPTIDES THROUGH EXPRESSION OF TITLE OF INVENTION: DNAS PLUE REPERENCE: 382.1026

FILE REFERENCE: 382.1026

CURRENT FILING DATE: 1999-03-26

EARLIER APPLICATION NUMBER: US/09/280,030A

CURRENT FILING DATE: 1999-03-26

EARLIER APPLICATION NUMBER: JP10-87339/1998

EARLIER FILING DATE: 1998-03-31

NUMBER OF SEQ ID NOS: 66

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 8
 SOFTWARE: PatentIn version 3.0
SEQ ID NO 666
LENGTH: 17
 Sequence 666, Applic Publication No. US20 GENERAL INFORMATION:
 Query Match
Best Local Similarity
Matches 15; Conserv
 SEQ ID NO 303
LENGTH: 17
 GENERAL INFORMATION:
 Sequence 8, Application US/09280030A Patent No. US20010021515A1
 Matches
 Query Match
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, Jim TITLE OF INVENTION: Nucleic acid treatment TITLE OF INVENTION: HER2 FILE REFERENCE: MBHB01-1653-A (400/014) CURRENT APPLICATION NUMBER: US/10/163,552 CURRENT FILING DATE: 2002-06-06 NUMBER OF SEQ ID NOS: 1997
 APPLICANT: Sato, Seiji
APPLICANT: Higashikuni, Naohiko
APPLICANT: Kudo, Toshiyuki
APPLICANT: Kondo, Masaaki
 OTHER INFORMATION: Description of Artificial OTHER INFORMATION: a forward oligonucleotide
 ORGANISM: Artificial Sequence FEATURE:
 LENGTH: 18
TYPE: DNA
 TYPE: RNA
 ENGTH:
 Local Similarity
les 12; Conserv
 2269 CCAGTCAAGTGGATGGC 2285
 1920 TCTTCTTGGAGCCTGCA 1936
 17 TCTTCTTGCAGCCAGCA 1
 1 CCCAUCAAGUGGAUGGC 17
 Application US/10163552
o. US20030105051A1
 Conservative
 Conservative
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 Score 13.8; D
Pred. No. 1.2e
3; Mismatches
 Score 13.8;
Pred. No. 1
 Mismatches
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1.2e+02;
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 DB 1;
 DB
 Sequence:
encoding (
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 Length 17;
 Indels
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Query Match

Score 13.8;

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Length 18

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RESULT 97
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 ; OTHER INFORMATION: Description of Artificial Sequence: Designated is ; OTHER INFORMATION: a reverse oligonucleotide encoding (His)6 US-09-280-030-9
 RESULT 96
US-09-280-030-9
 В
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 US-10-257-848-13
 GENERAL INFORMATION:
 Matches
 Query Match
Best Local
 Sequence 9, Application US/09280030A Patent No. US20010021515A1
 Best Local Similarity
 Matches
 APPLICANT:
APPLICANT:
 PRIOR FILING DATE: 2001-04-20
PRIOR APPLICATION NUMBER: JP 12-126340
PRIOR PILING DATE: 2000-04-21
PRIOR APPLICATION NUMBER: JP 12-205587
PRIOR FILING DATE: 2000-07-03
PRIOR APPLICATION NUMBER: JP 12-247962
PRIOR APPLICATION NUMBER: JP 12-247962
 TITLE OF INVENTION: DNAS ENCODING NEW FUSION PROTEINS AND PROCESSES FOR TITLE OF INVENTION: PREPARING USEFUL POLYPEPTIDES THROUGH EXPRESSION OF FILE REFERENCE: 382.1026
CURRENT APPLICATION NUMBER: US/09/280,030A
CURRENT FILING DATE: 1999-03-26
EARLIER FILING DATE: 1999-03-31
EARLIER FILING DATE: 1998-03-31
NUMBER: OF SEQ ID NOS: 66
 APPLICANT: Sato, Seiji
APPLICANT: Higashikuni, Nac
APPLICANT: Kudo, Toshiyuki
APPLICANT: Kondo, Masaaki
 CURRENT APPLICATION NUMBER: US/10/257,848
CURRENT FILING DATE: 2002-10-17
 APPLICANT: OGI, Kazuhiro
TITLE OF INVENTION: No. US20030158381A1el Insulin/IGF/Relaxin Family Polypeptide and
FILE REFERENCE: 2717 USOP
 APPLICANT: ITOH, APPLICANT: SUZU
 SOFTWARE:
PRIOR APPLICATION NUMBER: JP 12-395050 PRIOR FILING DATE: 2000-12-22
 CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: PCT/JP01/03399
 PRIOR FILING DATE: 2000-08-10
 LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
 FEATURE:
 Local
 1879 GAGATGATGAAGATGAT 1895
 1879 GAGATGATGAAGATGAT 1895
 1.0%;
1 Similarity 88.2%;
15; Conservative 0
 PatentIn Ver. 2.0
 GTGATGATGATGAT 17
 NISHI, Kazunori
KIZAWA, Hideki
HARADA, Masataka
 SUZUKI, No. US20030158381A1uhiro
 Application US/10257848
 Conservative
 Yasuaki
 88.2%;
 Naohiko
 0
 Pred. No. 1.4e+02;
0; Mismatches 2;
 Score 13.8; DB 1; Pred. No. 1.4e+02;
 Length 18
 Indels
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; OTHER INFORMATION: Primer US-10-257-848-13
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 US-10-004-551-43
 RESULT 99
US-10-004-551-43/c
 US-10-004-551-40
 RESULT 98
US-10-004-551-40
 SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 40
 GENERAL INFORMATION:
 SEQ ID NO 43
LENGTH: 18
 Sequence 40, Application US/10004551
Publication No. US20030004310A1
 Best Local Similarity 88.2%;
Matches 15; Conservative
 Best Local Similarity Matches 15; Conserv
 GENERAL INFORMATION
 Sequence 43, Application US/10004551 Publication No. US20030004310A1
 Matches
 Query Match
 Query Match
 Best Local Similarity
 APPLICANT: FERNANDES, ELMA
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES ENCODED THEREBY
FILE REFERENCE: 15966-559
CURRENT APPLICATION NUMBER: US/10/004,551
CURRENT FILING DATE: 2001-12-05
PRIOR APPLICATION NUMBER: 09/635,949
PRIOR APPLICATION NUMBER: 09/635,949
PRIOR FILING DATE: 2000-08-10
 PRIOR APPLICATION NUMBER: 09/635,949
PRIOR FILING DATE: 2000-08-10
NUMBER OF SEQ ID NOS: 110
 CURRENT APPLICATION NUMBER: US/10/004,551
CURRENT FILING DATE: 2001-12-05
 APPLICANT: SHIMKETS, RICHARD A
APPLICANT: FERNANDES, ELMA
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES
PILE REFERENCE: 15966-559
 APPLICANT: SHIMKETS, RICHARD A APPLICANT: FERNANDES, ELMA
 SOFTWARE: PatentIn Ver. 2.1
 NUMBER OF SEQ ID NOS: 110
 ORGANISM: Artificial Sequence
 TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
 OTHER INFORMATION: Description of Artificial Sequence: PCR PRIMER
 FEATURE:
 ORGANISM: Artificial Sequence
 TYPE:
 OTHER INFORMATION: Description of Artificial Sequence: PCR PRIMER
 FEATURE:
 TYPE: DNA
 LENGTH:
 LENGTH:
 1481 CGACCAAGAAGCCAGAC 1497
 1653 GGCAGGGGTCTCCGAGT
 15;
 18
 N
 2 CTACCAAGAAGCCAGCC
 GGCAGGGGTCTCTGTGT 18
 Conservative
 1.0%;
 1.0%;
 1669
 0;
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Score 13.8; DB 1;
Pred. No. 1.4e+02;
0; Mismatches 2
 Score 13.8; DB 1
Pred. No. 1.4e+02
 Score 13.8; DB 1;
Pred. No. 1.4e+02;
 Mismatches
 Mismatches
 DB 1;
 DB 1;
 Length 18;
 Length 18;
 Indels
 Indels
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 Gaps
 Gaps
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US-09-274-553D-999/c
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 RESULT 100
US-09-504-231A-999/c
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 US-09-504-231A-999
 CURRENT FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 09/274,53
PRIOR FILING DATE: 1999-03-23
PRIOR APPLICATION NUMBER: 09/257,608
PRIOR APPLICATION NUMBER: 09/257,608
PRIOR FILING DATE: 1998-02-24
PRIOR FILING DATE: 1998-09-18
PRIOR FILING DATE: 1998-09-18
 APPLICANT: PAYCO, PAMELA
APPLICANT: PAYCO, PAMELA
APPLICANT: MACEJAK, DENNIS
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES OR CONDITIONS RELATE
TITLE OF INVENTION: HEPATITIS C VIRUS INFECTION
FILE REFERENCE: rpi 247/282
CURRENT APPLICATION NUMBER: US/09/274,553D
CURRENT FILING DATE: 1999-03-23
PRIOR APPLICATION NUMBER: 09/257,608
PRIOR FILING DATE: 1999-02-24
PRIOR APPLICATION NUMBER: 60/100,842
PRIOR FILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 60/083,217
PRIOR FILING DATE: 1998-04-27
NUMBER OF SEQ ID NOS: 3148
 Sequence 999, Application US/09274553D Patent No. US20020082225A1
 Query Match
Best Local Similarity
 Matches
 SEQ ID NO 999
 GENERAL INFORMATION:
 Sequence 999, Application US/09504231A Patent No. US20020013458A1
 SOFTWARE: PatentIn version 3.0 EQ ID NO 999
 APPLICANT:
 APPLICANT: Macejak, Dennis
TITLE OF INVENTION: ENZYMATIC NUCLBIC ACID TREATMENT OF DISEASES OR CONDITIONS RELATE
TITLE OF INVENTION: HEPATITIS C VIRUS INFECTION
FILE REFERENCE: rpi 247/282
CURRENT APPLICATION NUMBER: US/09/504,231A
CURRENT FILING DATE: 2000-02-15
 SOFTWARE: PatentIn version 3.0
 PRIOR APPLICATION NUMBER: 60/083,217
PRIOR FILING DATE: 1998-04-27
 APPLICANT: Blatt, Lawrence
 NUMBER OF SEQ ID NOS: 3242
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICANT: Blact,
 FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:
 LENGTH: 15
TYPE: RNA
ORGANISM: Artificial Sequence
NO 999
TH: 15
 2433 CAGAATGGATAAGCC 2447
 1481 CGACCAAGAAGCCAGAC 1497
 INFORMATION:
 17
 14;
 McSwiggen, James
Roberts, Beth
 Roberts, Beth
 McSwiggen,
 Pavco,
 Conservative
 Lawrence
 Pamela
 1.0%;
 James
 Score 13.4; DB 1;
Pred. No. 1.1e+02;
 Mismatches
 .1e+02;
 Length 15;
 Nucleic
 <u>,</u>
 Acid
 Gaps
 Target
 0
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 ; TYPE: RNA
; OYANISM: Artificial Sequence
; FEATURE:
;, OTHER INFORMATION: Description of Artificial Sequence:
US-09-274-553D-999
 US-10-277-494-73
 RESULT 103
US-10-277-494-73
 US-10-277-494-3
 RESULT 102
US-10-277-494-3
 Ś
 SOFINAL 15
 TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or conditions Related To Level:
TITLE OF INVENTION: Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-K (400/064)
CURRENT APPLICATION NUMBER: US/10/277,494
CURRENT FILING DATE: 2002-10-21
NUMBER OF SEQ ID NOS: 446
SOFTWARE: PatentIn version 3.0
SEQ ID NO 73
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals,
APPLICANT: McSwiggen, Jim
 Matches
 Query Match
 GENERAL INFORMATION:
 Sequence 73, Application US/10277494 Publication No. US20030186909A1
 Sequence 3, Application US/10277494 Publication No. US20030186909A1
 Matches
 Best Local Similarity
 Query Match
 Best Local Similarity Matches 14; Conserv
 Query Match
 CURRENT APPLICATION NUMBER: US/10/277,494
CURRENT FILING DATE: 2002-10-21
NUMBER OF SEQ ID NOS: 446
 TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or conditions Related TITLE OF INVENTION: Epidermal Growth Factor Receptors FILE REFERENCE: MBHB00-958-K (400/064)
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, Jim
 LENGTH: 15
TYPE: RNA
ORGANISM: Homo sapiens
 LENGTH:
TYPE: RI
 SOFTWARE: PatentIn version 3.0
 TYPE: RNA
ORGANISM: Homo sapiens
 Local Similarity
 2317 CATCAGAGTGATGTC 2331
 2318 ATCAGAGTGATGTCT 2332
 2433 CAGAATGGATAAGCC 2447
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 ACCAGAGUGAUGUCU 15
 CAGAGTGGATAAGCC
 Conservative
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 Conservative
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 Score 13.4; DB 1;
Pred. No. 1.1e+02;
 Score 13.4; DB 1;
Pred. No. 1.1e+02;
 Score 13.4; DB 1;
Pred. No. 1.1e+02;
0; Mismatches 1;
 Mismatches
 Mismatches
 Length 15;
 Length 15;
 Length 15;
 Indels
 Indels
 Nucleic Acid Target
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 Gaps
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 Level
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 RESULT 105
US-09-866-108-8355/c
 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-8354
 US-09-866-108-8354/c
 RESULT 104
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G
APPLICANT: HANZEL, David K
APPLICANT: RANK, David R.
 Sequence 8355, Application US/09866108 Patent No. US20020048800A1
 Query Match
Best Local (
 SEQ ID NO 8354
LENGTH: 17
 GENERAL INFORMATION:
 Matches
 Patent No.
 equence 8354, Application US/09866108 atent No. US20020048800A1
 APPLICANT:
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 SOFTWARE: Aeomica Sequence Listing Engine
 PRIOR APPLICATION PRIOR FILING DATE:
 APPLICANT: SHANNON, MAXK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04 APPLICATION NUMBER: US 60/236,
 APPLICATION NUMBER: US 60/207,456 FILING DATE: 2000-05-26
 FILING DATE: 2000-09-27
 APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00666
 FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
 APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00667
 1573 TCCAGCTCCTCCATG 1587
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00670
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00668
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00663
 INFORMATION:
 ILING DATE:
 LING DATE:
 LING DATE: 2001-01-30
 [LING DATE: 2001-01-30
 ILING DATE:
 17 rccaccrccrccrrc 3
 LING DATE:
 14;
 Similarity
 PENN, Sharron G.
HANZEL, David K.
 JI, Yonggang
 Conservative
 AEOMICA-7
 Wensheng
 David R.
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 1.0%;
 US 60/236,359
 . ი
 Score 13.4; DB 1; Length 17; Pred. No. 1.4e+02;
 0;
 Gaps
 0
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뭥
 US-09-866-108-8356/c
 RESULT 106
 ; TYPE: DNA
; ORGANISM: Homo
US-09-866-108-8355
 GENERAL
 Sequence 8356, Application US/09866108 Patent No. US20020048800A1
 Matches
 Query Match
Best Local
 SOFTWARE: Aeo
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
 APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 PRIOR
PRIOR
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
 APPLICANT: GU, Yizhong
 PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR FILING DATE: 2000-05-26 PRIOR APPLICATION NUMBER: GB PRIOR FILING DATE: 2000-10-04
 FILE REFERENCE: AEOMICA-7
 NUMBER OF
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: US 60/236,359
 PRIOR APPLICATION NUMBER: PCT/US01/00666
 RIOR FILING DATE:
 ILE REFERENCE: AEOMICA-7
 .PPLICANT: CHEN, Wensheng
.PPLICANT: SHANNON, Mark
.TTLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 LENGTH:
 ocal Similarity
 APPLICATION NUMBER: US 60/266,860
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: US 60/234,687
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00662
 1573 TCCAGCTCCTCCATG 1587
 FILING DATE: 2000-09-21
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00670
 INFORMATION:
 16 TCCAGCTCCTCCTTG
 LING DATE:
 14;
 ING DATE: 2001-02-05
SEQ ID NOS: 15752
 Aeomica Sequence
 Conservative
 sapiens
 2001-01-30
 2001-01-30
 2001-01-30
 2000-09-27
 1.0%;
 GB 24263.6
 Listing Engine
 ٥,
 Score 13.4;
Pred. No. 1
 Mismatches
 DB 1;
..4e+02;
 Length 17;
 Indels
 0
 Gaps
 0
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US-09-866-108-8373/c
 片
 S
 ; SEQ ID NO 8356
; LENGTH: 17
; TYPE: DNA
; ORCANISM: Homo sapiens
US-09-866-108-8356
 APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 Sequence 8373, Application Patent No. US20020048800A1
 Query Match
Best Local :
 Matches
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggai
APPLICANT: PENN, Shar:
 PRIOR
PRIOR
 SOFTWARE: Aeomica Sequence
 PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
 OR APPLICATION NUMBER: US 60/236,359
OR FILING DATE: 2000-09-27
OR APPLICATION NUMBER: PCT/US01/00666
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00667
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00664
OR APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: GB 24263.6
FILING DATE: 2000-10-04
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: US 60/266,860
 APPLICATION NUMBER: US 60/234,687
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00669
 FILING DATE: 2000-09-
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00662
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00667
 APPLICATION NUMBER: PCT/US01/00666 FILING DATE: 2001-01-30
 1573 TCCAGCTCCTCCATG 1587
 FILING DATE: 2001-01-30
 FILING DATE:
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: US 60/236,359
 INFORMATION:
 15
 1.0%;
Similarity 93.3%;
14; Conservation
 Yonggang
 2001-01-30
 2000-09-
 US/09866108
 Listing Engine
 Score 13.4; DB 1;
Pred. No. 1.4e+02;
 ä
 HUMAN HEART
 Length 17;
 Indels
 <u>0</u>
 Gaps
 0
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 US-09-866-108-8373
 PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR PPLICATION NUMBER: US 60/236,359
PRIOR PRILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR PRILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
 US-09-866-108-8376/c
PRIOR FILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00665

PRIOR APPLICATION NUMBER: PCT/US01/00668

PRIOR APPLICATION NUMBER: PCT/US01/00668

PRIOR PILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00663

PRIOR FILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00662

PRIOR APPLICATION NUMBER: PCT/US01/00661

PRIOR APPLICATION NUMBER: PCT/US01/00661
 SOFTWARE: Aeoma
SOFTWARE: Aeoma
; SEQ ID NO 8373
 Best Loc
Matches
 Sequence 8376, Application US/09866108 Patent No. US20020048800A1
 GENERAL
 Query Match
 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
 APPLICANT: GU, Yizhong APPLICANT: JI, Yongga
 PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
 APPLICANT:
 PRIOR
 APPLICANT: SHANNON, Mark
IITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
 APPLICANT:
 NUMBER OF SEQ ID NOS: 15752
 TYPE: DNA
ORGANISM: Homo sapiens
 Local Similarity
nes 14; Conserv
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00665
 1507
 INFORMATION:
 17
 Aeomica Sequence Listing Engine
 CAGCCGGCTGTGCAC 1521
 PENN, Sharron G. HANZEL, David K.
 CHEN, Wensheng
 RANK, David R.
 Conservative
 1.0%;
 Score 13.4;
Pred. No. 1.
 4e+02;
 DB
 ,
--
 IN HUMAN
 Length 17;
 HEART
 AND MUSCLE
 0
 Gaps
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US-09-866-108-8943; Sequence 8943; Application US/09866108; Patent No. US20020048800A1
 밁
 ; ORGANISM: Homo sapiens
US-09-866-108-8376
 RESULT 109
 Matches
 Query Match
 PRIOR APPLICATION NUMBER: US 60/266,860 PRIOR FILING DATE: 2001-02-05 NUMBER OF SEQ ID NOS: 15752 SOFTWARE: Accession 2001-02-05 PRIOR PRI
 APPLICANT:
 SOFTWARE: Aeomica Sequence Listing Engine
 APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 CURRENT APPLICATION NUMBER: US/09/866,108
SOFTWARE: Aeomica Sequence Listing Engine
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: US 60/266,860
 TYPE: DNA
 ENGTH:
 ID NO 8376
 Local
 APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
 APPLICATION NUMBER: GB 24263.6
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00665 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00666
 APPLICATION NUMBER: US 60/236,359
 FILING DATE: 2000-10-04
 1506 CCAGCCGGCTGTGCA 1520
 APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00663
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00667
 APPLICATION NUMBER: US 60/234,687
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668
 FILING DATE:
 OF SEQ ID NOS: 15752
 INFORMATION
 15
 ILING DATE:
 LING DATE:
 LING DATE:
 LING DATE:
 14;
 LING DATE:
 Similarity
 CCAGCTGGCTGCA 1
 HANZEL,
 Conservative
 Yizhong
 Yonggang
NN, Sharron G.
NZEL, David K.
 AEOMICA-
 2001-02-05
 2000-09-21
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 2000-09-27
 1.0%;
 2001-05-25
 ٥,
 Score 13.4; DB 1;
Pred. No. 1.4e+02;
0; Mismatches 1;
 Length 17;
 Indels
 0;
 Gaps
 0
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; TYPE: DNA; ORGANISM: Homo sapiens US-09-866-108-8943
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 ; ORGANISM: Homo sapiens
US-09-866-108-8944
 RESULT 110
US-09-866-108-8944
 Sequence 8944, Appli
Patent No. US2002004
GENERAL INFORMATION:
 Query Match
Best Local Similarity
Matches 14; Conser
 SEQ ID NO 8944
 Matches
 Query Match
Best Local Similarity
 PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 SOFTWARE: Aeomica Sequence Listing Engine
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 NUMBER OF SEQ ID NOS: 15752
 PRIOR APPLICATION I
 PRIOR
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 PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
 APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
 APPLICANT: GU, Yizhong
 PRIOR APPLICATION NUMBER: PCT/US01/00670
 PRIOR APPLICATION NUMBER: PCT/US01/00665
 PRIOR FILING DATE: 2000-09-27
 PRIOR APPLICATION NUMBER: US 60/236,359
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: GB 24263.6
 APPLICANT:
 RIOR FILING DATE:
 RIOR APPLICATION NUMBER: PCT/US01/00662
 PPLICANT:
 LENGTH:
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: US 60/266,860 FILING DATE: 2001-02-05
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER:
 FILING DATE:
 FILING DATE: 2001-01-30
1355 CAGCGCCTGGAAGAG 1369
 FILING DATE: 2000-09-21
 FILING DATE: 2001-01-30
 1355 CAGCGCCTGGAAGAG 1369
 14;
 3 CGGCGCCTGGAAGAG 17
 CATION NUMBER: US 60/234,687
 JI, Yonggang
PENN, Sharron
 RANK, David R. CHEN, Wensheng
 HANZEL, David
 Application
0020048800A1
 Conservative
 Conservative
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 2000-10-04
 1.0%;
 1.0%;
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 PCT/US01/00663
 US/09866108
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 Score 13.4; I
Pred. No. 1.4e
0; Mismatches
 Score 13.4;
Pred. No. 1
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 1.4e+02;
 .4e+02;
 DB 1;
 DB
 IN HUMAN HEART AND
 Length 17;
 Length 17;
 Indels
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 0,
 0
 MUSCLE
 Gaps
 Gaps
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 0
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CGGCGCCTGGAAGAG 16

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RESULT 112
US-09-866-108-9000
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 ; ORGANISM: Homo sapiens
US-09-866-108-8945
 US-09-866-108-8945
GENERAL INFORMATION:
 Sequence 9000, Application US/09866108 Patent No. US20020048800A1
 Query Match
Best Local Similarity
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 NUMBER OF SEQ ID NOS: 15752
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
 APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
 SOFTWARE: Acomica Sequence Listing Engine
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT:
 TYPE: DNA
 ENGTH:
 ID NO 8945
 FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
 APPLICATION NUMBER: US 60/236,359 FILING DATE: 2000-09-27
 ce 8945, Application US/09866108
 1355 CAGCGCCTGGAAGAG 1369
 APPLICATION NUMBER: US 60/234,687
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00663
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
 INFORMATI
 LING DATE: 2001-01-30
 14;
 LING DATE: 2001-01-30
 1 CGGCGCCTGGAAGAG 15
 RANK, David R. CHEN, Wensheng
 HANZEL,
 PENN, Sharron
 Conservative
 Yonggang
 AEOMICA-7
 1.0%;
 2001-05-25
 ო ი
 Score 13.4;
Pred. No. 1.
 .4e+02
 IN HUMAN HEART AND MUSCLE
 Length 17;
 Indels
 0,
 Gaps
 0
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RESULT 113
US-09-866-108-9003
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 Ś
 ; ORGANISM: Homo sapiens US-09-866-108-9000
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharro
APPLICANT: HANZEL, Dav
 GENERAL
 Sequence 9003, Application US/09866108 Patent No. US20020048800A1
 Query Match
Best Local Similarity
Matches 14; Conserv
 SOFTWARE: Aeo
 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
 PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
 NUMBER OF SEQ ID NOS: 15752
 PRIOR
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00668 PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
 PRIOR APPLICATION NUMBER: PCT/US01/00669
 PRIOR APPLICATION NUMBER: PCT/US01/00661
 PRIOR APPLICATION NUMBER: PCT/US01/00663
 FILE REFERENCE:
 LENGTH: 17
 PPLICANT: SHANNON, Mark
THE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662 FILING DATE: 2001-01-30
 FILING DATE: 2
 1505 GCCAGCCGGCTGTGC 1519
 INFORMATION:
 w
 Aeomica Sequence Listing Engine
 GCCAGCCGGCGGTGC 17
 PENN, Sharron (
HANZEL, David F
RANK, David R.
 CHEN, Wensheng
 Conservative
 AEOMICA-7
 001-01-30
 1.0%;
 0;
 Score 13.4;
Pred. No. 1.
 Mismatches
 4e+02
 DB
 1;
 IN HUMAN HEART AND MUSCLE
 Length 17;
 Indels
 0,
 Gaps
 0,
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밁
 á
 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-9003
 US-09-825-805-648
 RESULT 114
 Sequence 648, Application US/09825805
Publication No. US20030004122A1
 SEQ ID NO 9003
 Matches
 Best Local
 Query Match
 GENERAL INFORMATION:
 APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleotide Triphosphate and their Incorporation into Oligonucleot
FILE REFERENCE: MBHB00-831-F (400/009)
CURRENT APPLICATION NUMBER: US/09/825,805
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: 09/578,223
PRIOR APPLICATION NUMBER: 09/578,223
PRIOR FILING DATE: 2000-05-23
PRIOR FILING DATE: 2000-05-23
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Beigelman, Leo
 NUMBER OF SEQ ID NOS: 15752
 PRIOR FILING DATE: 1999-12-30
PRIOR APPLICATION NUMBER: 09/474,432
 PRIOR APPLICATION NUMBER: 09/476,387
 APPLICANT:
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 SOFTWARE: Aeomica Sequence Listing Engine
 PRIOR
 APPLICANT:
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 ENGTH:
 FILING DATE: 1999-12-29
APPLICATION NUMBER: 09/301,511
FILING DATE: 1999-04-28
 APPLICATION NUMBER: US 60/234,687
 1506 CCAGCCGGCTGTGCA 1520
 FILING DATE:
 APPLICATION NUMBER: US 60/266,860
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00665
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00667
 FILING DATE:
 APPLICATION NUMBER: US 60/236,359
 APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04
 FILING DATE:
 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
 FILING DATE: 2001-01-30
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00666
APPLICATION NUMBER: 09/186,675
 APPLICATION NUMBER: PCT/US01/00670
 ILING DATE:
 ILING DATE:
 14:
 Similarity
 CCAGCCGGCGGTGCA 15
 Sweedler, Dave
 Adamic, Jasenka Matulic
 Karpeisky, Alex
 Beaudry, Amber
 Conservative
 2000-05-26
 2001-01-30
 2000-09-27
 2001-02-05
 2000-09-
 2001-01-30
 2001-01-30
 2001-01-30
 Dave
 1.0%;
 <u>.</u>
 Score 13.4; DB 1; Length 17; Pred. No. 1.4e+02; o; Mismatches 1; Indels
 <u>,</u>
 0
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 ; TYPE: RNA; Homo sapiens US-09-780-533A-2375
 밁
 S
 ; TYPE: RNA; Homo sapiens US-09-825-805-648
 US-09-780-533A-2376/c
 US-09-780-533A-2375/c
 RESULT 115
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
 GENERAL INFORMATION:
 SOFTWARE: Par
SEQ ID NO 648
SEQ ID NO 2376
LENGTH: 17
 Sequence 2376, Application US/09780533A Publication No. US20030060611A1
 Matches
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 SEQ ID NO 2375
 Sequence 2375, Application US/09780533A Publication No. US20030060611A1
 Query Match
Best Local Similarity
Matches 11; Conserv
 APPLICANT: McSwiggen, Jim
APPLICANT: Chowrira, Bharat
APPLICANT: Haeberii, Pete
APPLICANT: Haeberii, Pete
FIIER OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
CURRENT FILING DATE: 2001-02-11
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
 PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: Patentin version 3.0
 PRIOR APPLICATION NUMBER: US 60/181,797 PRIOR FILING DATE: 2000-02-11
 APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
 PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-11-05
 CURRENT APPLICATION NUMBER: US/09/780,533A CURRENT FILING DATE: 2001-02-09
 TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene FILE REFERENCE: MBHB00,878-A (400/011)
 NUMBER OF SEQ ID NOS: 1558
 SOFTWARE: PatentIn version 3.0
 APPLICANT:
 APPLICANT:
 LENGTH:
 LENGTH: 17
 Local Similarity
 1573 TCCAGCTCCTCCATG 1587
 2110 GGCATGGAGTACTTG 2124
 17 TCCAGCTCCTCCAGG
 14;
 17
 1 GGCAUGGAGCACUUG 15
 PatentIn version 3.0
 McSwiggen, Jim
Chowrira, Bharat
Haeberli, Pete
 Conservative
 Conservative
 1.0%;
73.3%;
 93.3%;
 1.0%;
 60/064,866
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 0,
 Score 13.4; DB 1;
Pred. No. 1.4e+02;
 Score 13.4; DB 1;
Pred. No. 1.4e+02;
 Mismatches
 Mismatches
 Length 17;
 Length 17;
 Indels
 0
 0
 Gaps
 Gaps
 0
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US-09-877-478-2527
 밁
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 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-780-533A-2376
 ; LENGTH: 17
; TYPE: RNA
; ORGANISM: Hepatitis B virus
US-09-877-478-2182
 밁
 Ś
 NUMBER OF SEQ ID NOS: 6586
SOFTWARE: PatentIn version 3.0
SEQ ID NO 2182
 GENERAL INFORMATION:
 Sequence 2527, Application US/09877478 Publication No. US20030068301A1
 1.0%;
Best Local Similarity 93.3%;
Matches 14; Conservative
 Query Match
Best Local
 Matches
 GENERAL INFORMATION:
APPLICANT: McSwiggen, Jim
APPLICANT: McTrilsey, Dave
TITLE OF INVENTION: Method and Reagent for Inhibiting Hepatitis B Virus Replication
FILE REFERENCE: MBHB00-845-H (400/029)
CURRENT APPLICATION NUMBER: US/09/877,478
 APPLICANT: Ribozyme Pharmaceuticals, APPLICANT: Draper, Kenneth APPLICANT: Blatt, Larry
 APPLICANT: McSwiggen, Jim
APPLICANT: McKrissey, Dave
TITLE OF INVENTION: Method and Reagent for Inhibiting Hepatitis B Virus Replication
FILE REFERENCE: MSHB00-845-H (400/029)
CURRENT APPLICATION NUMBER: US/09/877,478
CURRENT FILING DATE: 2001-12-31
 PRIOR APPLICATION NUMBER: US 09/636,385
PRIOR FILING DATE: 2000-08-09
PRIOR APPLICATION NUMBER: US 09/696,347
PRIOR FILING DATE: 2000-10-24
PRIOR APPLICATION NUMBER: US 08/193,627
PRIOR FILING DATE: 1994-02-07
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Draper, Kenneth APPLICANT: Blatt, Larry
 PRIOR APPLICATION NUMBER: US 09/436,430 PRIOR FILING DATE: 1999-11-08
 PRIOR FILING DATE: 1992-05-14
PRIOR APPLICATION NUMBER: US 09/531,025
 PRIOR APPLICATION NUMBER: US 07/882,712
 APPLICATION NUMBER: US 08/433,993
FILING DATE: 1995-05-04
APPLICATION NUMBER: US 08/434,504
FILING DATE: 1995-05-04
 FILING DATE: 2000-03-20
 2479 ATGAGGGACTGTTGG 2493
 1573 TCCAGCTCCTCCATG 1587
 16
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 Similarity
 Draper, Kenn
Blatt, Larry
 AUGGGGGACUGUUGG 16
 TCCAGCTCCTCCAGG 2
 , Application US/09877478
o. US20030068301A1
 Conservative
 1.0%;
 <u>,,</u>
 Score 13.4;
Pred. No. 1.
 Score 13.4; DB 1;
Pred. No. 1.4e+02;
 Mismatches
 Mismatches
 1.4e+02;
 DB 1;
 Length 17;
 Length 17;
 Indels
 0
 0
 0,
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뭉
 Query Match
Best Local Similarity
Watches 14; Conserve
 ; TYPE: RNA; Homo sapiens US-09-792-818-96
 RESULT 119
US-09-792-818-96/c
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 ; ORGANISM: Hepatitis B virus US-09-877-478-2527
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 96 LENGTH: 17
 Sequence 96, Application US/09792818 Publication No. US20030134806A1
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 2527
 GENERAL INFORMATION:
 Matches
 Query Match
 CURRENT APPLICATION NUMBER: US/09/792,818
CURRENT FILING DATE: 2001-02-23
NUMBER OF SEQ ID NOS: 2304
 APPLICANT: Hamblin, Paul
APPLICANT: Ellis, Jonathan
TITLE OF INVENTION: Method and Reagent for
TITLE OF INVENTION: (GRID) Gene
FILE REFERENCE: MBHB00-901-A (400/013)
 APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Jarvis, Thale
APPLICANT: Von Carlowitz, Ira
 APPLICANT:
 CURRENT FILING DATE: 2001-12-31
PRIOR APPLICATION NUMBER: US 07/882,712
PRIOR FILING DATE: 1992-05-14
PRIOR APPLICATION NUMBER: US 09/531,025
 PRIOR APPLICATION NUMBER: US 09/436,430 PRIOR FILING DATE: 1999-11-08
 PRIOR FILING DATE: 1994-02-07
PRIOR APPLICATION NUMBER: US 08/433,993
PRIOR FILING DATE: 1995-05-04
 PRIOR APPLICATION NUMBER: US 09/696,347
PRIOR FILING DATE: 2000-10-24
PRIOR APPLICATION NUMBER: US 08/193,627
 NUMBER OF SEQ ID NOS: 6586
 PRIOR APPLICATION NUMBER: US 08/434,504
PRIOR FILING DATE: 1995-05-04
 PRIOR FILING DATE: 2000-03-20
PRIOR APPLICATION NUMBER: US 09/636,385
PRIOR FILING DATE: 2000-08-09
 Local Similarity
les 10; Conserv
 1868 TGTCAGAGATGGAGA 1882
 2479 ATGAGGGACTGTTGG 2493
16 TGACAGAGATGGAGA 2
 AUGGGGGACUGUUGG 17
 Von Carlowitz, Ira
McSwiggen, Jim
 Conservative
 Conservative
 1.0%;
93.3%;
 1.0%; Score 13.4; DB 1; 66.7%; Pred. No. 1.4e+02;
 0
 Score 13.4; D
Pred. No. 1.4e
0; Mismatches
 Mismatches
 1.4e+02;
 the Inhibition of Grb-2-related with Inser
 DB 1;
 Length 17;
 Length 17;
 Indels
 0
 0,
 Gaps
 0
 0
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; Sequence 284, Application US/09792818 ; Publication No. US20030134806A1 ; GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc

RESULT 120 US-09-792-818-284/c

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US-10-230-006-164
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 FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
PRIOR APPLICATION NUMBER: PCT/US 02/16840
PRIOR FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: US 60/318,471
PRIOR FILING DATE: 2001-09-10
NUMBER OF SEQ ID NOS: 4666
 밁
 Ś
 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-238-700-3194
 US-10-238-700-3194/c
 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-792-818-284
 Sequence 164, Application US/10230006 Publication No. US20030191077A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity 93.3%;
Conservative
 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 284
 Query Match
 APPLICANT: Fosnaugh, Kathy
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: METHOD AND REAGENT FOR THE TREATMENT OF ASTHMA AND ALLERGIC CONDIFILE REFERENCE: 400/056 (MBHB01-1110)
CURRENT APPLICATION NUMBER: US/10/230,006
CURRENT FILING DATE: 2002-11-18
 TITLE OF INVENTION: Method and Reagent for the Inhibition of Grb-2-related with TITLE OF INVENTION: (GRID) Gene FILE REFERENCE: MBHB00-901-A (400/013) CURRENT APPLICATION NUMBER: US/09/792,818 CURRENT FILING DATE: 2001-02-23
 APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment
FILE REFERENCE: 400/057 (MBHB01-1158-A)
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, James
 NUMBER OF SEQ ID NOS: 2304
 APPLICANT: Ribozyme Pharmaceuticals, Inc
 SOFTWARE: PatentIn version 3.0
 APPLICANT:
 ENGTH:
 Local Similarity
APPLICATION NUMBER: US 60/315,315
 2283 GGCTCCAGAAGCCCT 2297
 1868 TGTCAGAGATGGAGA 1882
 17 GGCTCCAGCAGCCCT 3
 15 TGACAGAGATGGAGA 1
 Ellis, Jonathan
 Hamblin,
 McSwiggen, J
 Von Carlowitz,
 Jarvis,
 , Application US/10238700 o. US20030153521A1
 1.0%;
llarity 93.3%;
Conservative.
 1.0%; Score 13.4;
93.3%; Pred. No. 1
 Ira
 Score 13.4; DB 1;
Pred. No. 1.4e+02;
 Mismatches
 DB 1; Length 17;
 of Diseases or Conditions Related
 Length 17;
 ٥,
 0,
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망
 ; TYPE: RNA; Homo sapiens US-10-230-006-2204
 밁
 Query Match
Best Local Similarity
Matches 13; Conserva
 TYPE: RNA
; ORGANISM: Homo sapiens
US-10-230-006-164
 ; ORGANISM: Homo sapiens
US-10-230-006-2205
 US-10-230-006-2205/c
 RESULT 124
 RESULT 123
US-10-230-006-2204/c
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Fosnaugh, Kathy
APPLICANT: McSwiggen, Jim
 SEQ ID NO 2204
LENGTH: 17
 GENERAL INFORMATION:
 Sequence 2205, Application US/10230006 Publication No. US20030191077A1
 SOFTWARE: PatentIn version SEQ ID NO 164
 SEQ ID NO 2205
 Matches 14; Conserv
 Query Match
 Sequence 2204, Application US/10230006 Publication No. US20030191077A1
 Query Match
 APPLICANT: Fosnaugh, Kathy
APPLICANT: McSwiggen, Jim
IITLE OF INVENTION: METHOD AND REAGENT FOR THE TREATMENT OF ASTHMA AND ALLERGIC CONDI
FILE REFERENCE: 400/056 (MBHB01-1110)
CURRENT APPLICATION NUMBER: US/10/230,006
CURRENT FILING DATE: 2002-11-18
PRIOR APPLICATION NUMBER: US 60/315,315
PRIOR FILING DATE: 2001-08-28
NUMBER OF SEQ ID NOS: 2678
 APPLICANT: Fosnaugh, Kathy
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: METHOD AND REAGENT FOR
FILE REFERENCE: 400/056 (MBHB01-1110)
CURRENT APPLICATION NUMBER: US/10/230,006
CURRENT FILING DATE: 2002-11-18
PRIOR APPLICATION NUMBER: US 60/315,315
PRIOR FILING DATE: 2001-08-28
 NUMBER OF SEQ ID NOS: 2678
 APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: Fosnaugh, Kathy
 LENGTH: 17
TYPE: RNA
 SOFTWARE: PatentIn version 3.0
 SOFTWARE: PatentIn version 3.0
 NUMBER OF SEQ ID NOS: 2678
1.0%;
Local Similarity 93.3%;
es 14; Conservative
 1573 TCCAGCTCCATG 1587
 1993 GAATACCTCCGAGCC 2007
 17 TCCAGCTCCTCCAGG
 1 GAAGACCUCCGAGCC 15
 Conservative
 1.0%;
llarity 93.3%;
Conservative
 1.0%;
 <u>,</u>
 Score 13.4;
Pred. No. 1
 Score 13.4;
Pred. No. 1.
 Score 13.4;
Pred. No. 1.
 Mismatches
 Mismatches
 1.4e+02;
 .4e+02;
 .4e+02
 THE TREATMENT OF ASTHMA AND ALLERGIC CONDI
 DB 1;
 Length 17;
 Length 17;
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 0
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Matches

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1573

TCCAGCTCCTCCATG 1587

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RESULT 126
US-10-027-632-58632/c
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 ; TYPE: DNA
; ORGANISM: Human
US-10-027-632-58632
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 US-10-027-632-58632/c
 Sequence 58632,
 Matches
 Query Match
 SEQ ID NO 58632
 Sequence 58632, Application US/10027632 Publication No. US20030204075A9
 GENERAL INFORMATION:
 PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
 APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
 SENERAL INFORMATION:
 CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
 SOFTWARE: FastSEQ for Windows Version 4.0
 PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
 PRIOR APPLICATION NUMBER: US 60/218,006 PRIOR FILING DATE: 2000-07-12
 CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
 TITLE OF INVENTION: Identification and Mapping of Single Nucleotide TITLE OF INVENTION: Polymorphisms in the Human Genome FILE REFERENCE: 108827.129
 NUMBER OF SEQ ID NOS: 325720
 PRIOR APPLICATION NUMBER: US 60/198,676
 ENGTH: 17
 Local Similarity
 APPLICATION NUMBER: US 60/193,483
FILING DATE: 2000-03-29
APPLICATION NUMBER: US 60/185,218
FILING DATE: 2000-02-24
 APPLICATION NUMBER: US 60/198,676 FILING DATE: 2000-04-20
 APPLICATION NUMBER: US 60/167,363
FILING DATE: 1999-11-23
APPLICATION NUMBER: US 60/156,358
 APPLICATION NUMBER: US 60/146,002 FILING DATE: 1999-08-09
 FILING DATE: 2000-03-29
APPLICATION NUMBER: US 60/185,218
FILING DATE: 2000-02-24
 FILING DATE: 1999-08-09
 APPLICATION NUMBER: US 60/146,002
 FILING DATE: 1999-09-
 2326 GATGTCTGGTCCTTC 2340
 FILING DATE: 1999-09-28
 APPLICATION NUMBER: US 60/156,358
 APPLICATION NUMBER: US 60/167,363
 FILING DATE: 1999-11-
 15
 Conservative
 Application US/10027632
NOS:
 David G.
 1.0%;
 2002-04-30
 -23
 <u>,</u>
 Score 13.4; DB 1;
Pred. No. 1.4e+02;
 1.4e+02;
 Length 17;
 Indels
 <u>,</u>
 Gaps
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 RESULT 128
US-10-027-632-58640/c
 ; ORGANISM: Human US-10-027-632-58640
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 ; TYPE: DNA
; ORGANISM: Human
US-10-027-632-58632
 RESULT 127
 US-10-027-632-58640/c
 Sequence 58640, Application US/10027632
GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
ETTLE DEFENDED. 10000721
 SEQ ID NO 58640
 Matches
 Query Match
 APPLICANT: Wang, Da
 SEQ ID NO 58632
 Matches
 Sequence 58640, Application US/10027632 Publication No. US20030204075A9
 Query Match
Best Local Similarity
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
 TITLE OF INVENTION: Identification and Mapping of Single Nucleotide TITLE OF INVENTION: Polymorphisms in the Human Genome FILE REFERENCE: 100827.129
 PRIOR APPLICATION NUMBER: US 60/146,002 PRIOR FILING DATE: 1999-08-09
 PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
 CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
 SOFTWARE: FastSEQ for Windows Version 4.0
 SOFTWARE: FastSEQ for Windows Version
 PRIOR APPLICATION NUMBER: US 60/167,363 PRIOR FILING DATE: 1999-11-23
 PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
 PRIOR APPLICATION NUMBER: US 60/193,483
 TYPE: DNA
 PRIOR FILING DATE: 2000-02-24
 ENGTH: 17
 Local Similarity
les 14; Conserv
 2326 GATGTCTGGTCCTTC 2340
 FILING DATE: 1999-09-28
 APPLICATION NUMBER: US 60/156,358
 2326 GATGTCTGGTCCTTC 2340
 15
 14;
 15 GATGTCTGTTCCTTC 1
 Conservative
 ID NOS:
 Conservative
 1999-08-09
 1.0%;
 1.0%;
 0;
 Score 13.4; D:
Pred. No. 1.4e
0; Mismatches
 Score 13.4;
Pred. No. 1.
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 1.4e+02;
 .4e+02;
 DB 1;
 DB 1;
 Length 17;
 Length 17;
 Indels
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 0,
 0
 Gaps
 Gaps
 0,
 0
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PRIOR APPLICATION NUMBER: US 60/198,676

PPLICATION NUMBER: US 60/193,483

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 US-10-163-552-667
Sequence 667, Application US/10163552
Publication No. US20030105051A1
GENERAL INFORMATION:
 RESULT 129
US-10-163-552-289
 US-10-027-632-58640
 Publication No.
 SEQ ID NO 58640
 Best Local
 SEQ ID NO 289
 SEQ ID NO 667
 Query Match
 GENERAL INFORMATION:
 Matches 14; Conservative
 Best Local Similarity
 Query Match
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic acid treatment of diseases or conditions related to level
TITLE OF INVENTION: HER2
FILE REFERENCE: MBHB01-1653-A (400/014)
CURRENT APPLICATION NUMBER: US/10/163,552
CURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 1997
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, Jim TITLE OF INVENTION: Nucleic acid treatment TITLE OF INVENTION: HER2 FILE REFERENCE: MHHB01-1653-A (400/014) CURRENT EPLICATION UNUBER: US/10/163,552 CURRENT FILING DATE: 2002-06-06
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 PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
 SOFTWARE: FastSEQ for Windows Version 4.0
 NUMBER OF SEQ ID NOS: 1997
SOFTWARE: PatentIn version 3.0
 SOFTWARE: PatentIn version 3.0
 PRIOR
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 TYPE: DNA
ORGANISM: Human
ORGANISM: Homo'sapiens
 LENGTH: 17
TYPE: RNA
 ENGTH: 17
 FILING DATE: 2000-03-29
APPLICATION NUMBER: US 60/185,218
FILING DATE: 2000-02-24
APPLICATION NUMBER: US 60/167,363
 FILING DATE: 1999-11-23
APPLICATION NUMBER: US 60/156,358
FILING DATE: 1999-09-28
 2110 GGCATGGAGTACTTG 2124
 2326 GATGTCTGGTCCTTC 2340
 289,
 15 GATGTCTGTTCCTTC 1
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 Similarity
 Application US/10163552
 Conservative
 US20030105051A1
 1.0%;
 1.0%; Score 13.4; DB 1; Length 17; 73.3%; Pred. No. 1.4e+02;
 ٥,
 Score 13.4; DB Pred. No. 1.4e+(
 Mismatches
 Mismatches
 1.4e+02
 õ
 diseases
 Length 17
 conditions related
 0
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 Gaps
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APPLICANT: McSwiggen, Jim
ITILE OF INVENTION: Nucleic acid treatment of
ITILE OF INVENTION: HER2
ITILE OF INVENTION: MGHEN: US/10/163,552
ICURRENT APPLICATION NUMBER: US/10/163,552
ICURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 1997
INMBER OF SEQ ID NOS: 1997
INMBER OF SEQ ID NOS: 1997
INMBER OF SEQ ID NO 678
ILENCTH: 17
ITYPE: RNA
INGRANISM: Homo sapiens
US-10-163-552-678
 Query Match
Best Local Similarity
 , OTHER INFORMATION: Description of Artificial Sequence: US-09-848-754A-9124
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 RESULT 132
US-09-848-754A-9124
 RESULT 131
US-10-163-552-678
 US-10-163-552-667
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate.
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-1 (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
SQFTWARE: Patentin version 3.0
SSQ ID NO 9124
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: McSwiggen, Jim
 Matches
 Sequence 9124, Application US/09848754A Publication No. US20030073207A1
 Best Local Similarity Matches 10; Conserv
 Sequence 678, A
Publication No.
 Best Local Similarity Matches 11; Conserv
 Query Match
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 TYPE: RNA
ORGANISM: Artificial Sequence
 FEATURE:
 LENGTH: 13
 2273 TCAAGTGGATGGC 2285
 2320 CAGAGTGATGTCTGG 2334
 2273 TCAAGTGGATGGCTC 2287
 2 CAGAGUGAUGUGUGG 16
 1 UCAAGUGGAUGGCGC 15
 Application US/10163552
 Conservative
 Conservative
 Conservative
 US20030105051A1
 0.9%;
 1.0%;
 73.3%;
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 Score 13.4; DB 1;
Pred. No. 1.4e+02;
 Score 13.4;
Pred. No. 1
 Score 13; DB
Pred. No. 97;
 Mismatches
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 .4e+02;
 DB 1;
 DB 1;
 1;
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 Length 13;
 Length 17;
 Length 17;
 Indels
 Enzymatic Nucleic acid
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 0
 Gaps
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RESULT 133 US-10-277-494-54

Sequence 54, Application US/10277494

APPLICATION NUMBER: US 60/207,456

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RESULT 135
US-09-866-108-6887/c
 뭐
 Query Match
Best Local Similarity
Tatches 9; Conserve
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 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-277-494-54
 US-10-091-281-150/c
 RESULT 134
 US-10-091-281-150
 APPLICANT: SI ERWIN
APPLICANT: MORISETTE, JEAN
APPLICANT: MORISETTE, JEAN
TITLE OF INVENTION: OPTINEURIN NUCLEIC ACID MOLECULES AND USES
FILE REFERENCE: 13587.338
CURRENT APPLICATION NUMBER: US/10/091,281
CURRENT FILING DATE: 2002-03-06
NUMBER OF SEQ ID NOS: 463
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 150
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 Sequence 150, Application US/10091281 Publication No. US20030190617A1
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 54
 GENERAL INFORMATION:
 Patent No.
 Matches
 Best Local Similarity
 Query Match
 GENERAL INFORMATION:
 GENERAL INFORMATION:
 Publication No. US20030186909A1
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or
TITLE OF INVENTION: Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-K (400/064)
CURRENT APPLICATION NUMBER: US/10/277,494
CURRENT FILLING DATE: 2002-10-21
 APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSKIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga
 APPLICANT: RAYMOND, VINCENT
 NUMBER OF SEQ ID NOS: 446
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, Jim
 LENGTH: 16
TYPE: DNA
ORGANISM: Homo sapiens
 OTHER INFORMATION: Putative LYMF/TH1E47.01 motif
 FEATURE:
 ENGTH: 15
 2668 TCTCCAGACCCCA 2680
 2322 GAGTGATGTCTGG 2334
APPLICATION NUMBER: US/09/866,108 FILING DATE: 2001-05-25
 6887, Application o. US20020048800A1
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 GAGUGAUGUCUGG 13
 JI, Yonggang
 Conservative (
 Conservative
 0.9%;
 0.9%;
 US/09866108
 Score 13; DB 1; Le
s; Pred. No. 1.4e+02;
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 Score 13;
Pred. No.
 Mismatches
 DB 1;
1.3e+02;
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 Length 16;
 Length 15;
 HUMAN HEART
 Indels
 conditions Related
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 Gaps
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APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
ERIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
 RESULT 136
US-09-866-108-6892/c
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 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-6887
 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 6887
 Sequence 6892, Application US/09866108 Patent No. US20020048800A1
 Matches
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
 APPLICANT:
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 APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
 APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00661
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
 1359 GCCTGGAAGAGAA 1371
 APPLICATION NUMBER: PCT/US01/00662 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
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 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
 APPLICATION NUMBER: PCT/US01/00666
 FILING DATE:
 APPLICATION NUMBER: US 60/236,359
 FILING DATE: 2000-05-26
APPLICATION NUMBER: GB 24263.6
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 JI, Yonggang
PENN, Sharron G.
HANZEL, David K.
RANK, David R.
CHEN, Wensheng
 Conservative
 2001-01-30
 2000-09-27
 0.9%; Score 13; DB 1; L
100.0%; Pred. No. 1.6e+02;
 0
 Mismatches
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 Length 17;
 HUMAN HEART
 AND MUSCLE
 0,
 Gaps
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 RESULT 137
US-09-866-108-8946
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 SOFTWARE: Aeomica
SEQ ID NO 6892
 Query Match
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 Patent No.
 APPLICANT:
 PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
 CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 CURRENT APPLICATION NUMBER: US/09/866,108
 APPLICANT:
 LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
 PRIOR FILING DATE: 2001-02-05
 APPLICANT:
 PPLICANT:
 PPLICANT:
 PPLICANT: SHANNON, Mark .
ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 / Match 0.9%; So Local Similarity 100.0%; I nes 13; Conservative 0;
 FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00667
 APPLICATION NUMBER: US 60/236,359
 APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04
 APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
 APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663 FILING DATE: 2001-01-30
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00665 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00664
 1358 ССССТССААСАСА 1370
 APPLICATION NUMBER: US 60/266,860
 APPLICATION NUMBER: PCT/US01/00670
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00664
 FILING DATE:
 FILING DATE:
 8946, Application US/09866108
o. US20020048800A1
 13
 PENN, Sharron G. HANZEL, David K.
 RANK, David R
 Yizhong
 Yonggang
 AEOMICA-7
 Wensheng
NUMBER: PCT/US01/00663
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 2001-01-30
 2001-01-30
 2001-01-30
 Score 13; pred. No
 No.
 DB 1;
 1.6e+02;
 Length 17;
 0
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 ; ORGANISM: Homo sapiens US-09-866-108-8946
 GENERAL
 Sequence 8947, Application US/09866108 Patent No. US20020048800A1
 Query Match
 Matches
 PRIOR APPLIC
 PRIOR
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 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
 PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
 PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
 SOFTWARE:
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: PCT/US01/00670 PRIOR FILING DATE: 2001-01-30
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 PRIOR APPLICATION NUMBER: US 60/236,359 PRIOR FILING DATE: 2000-09-27
 APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 APPLICANT:
 APPLICANT:
 APPLICANT: GU, Yizhong
 NUMBER OF SEQ ID NOS: 15752
 PRIOR
 PRIOR
 PRIOR APPLICATION NUMBER: PCT/US01/00665
 PRIOR
 PRIOR FILING DATE: 2000-10-04
 APPLICANT:
 APPLICANT:
 APPLICANT:
 TYPE: DNA
 ID NO 8946
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 FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
 APPLICATION NUMBER: PCT/US01/00661
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00667 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00664
 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668
 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
 1357 GCGCCTGGAAGAG 1369
 INFORMATION
 2 GCGCCTGGAAGAG 14
 Aeomica Sequence Listing Engine
 CATION NUMBER: PCT/US01/00666
CATION NUMBER: US 60/266,860
 PENN, Sharron HANZEL, David
 CHEN, Wensheng
 RANK, David R
 0.9%; So larity 100.0%; I Conservative 0;
 Yonggang
 2001-02-05
 <u>ب</u> د
 Score 13; pred. No.
 Mismatches
 DB 1;
 1.6e+02;
 Length 17;
 Indels
 0;
 Gaps
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; ORGANISM: Homo sapiens US-09-866-108-9030
 RESULT 139
US-09-866-108-9030
 뭐
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 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-8947
 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 9030
Query Match
Best Local Similarity
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Best Local Similarity 100.
Matches 13; Conservative
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LENGTH: 17
 Patent No.
 NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engine
 LENGTH: 17
TYPE: DNA
 NUMBER OF SEQ ID NOS: 15752
 PRIOR APPLICATION NUMBER: UPRIOR FILING DATE: 2000-05-
 CURRENT APPLICATION NUMBER: US/09/866,108
 FILE REFERENCE: AEOMICA-
 RITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
 APPLICANT: GU, Yizhong
 OR APPLICATION NUMBER: PCT/US01/00667
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00664
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00669
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: US 60/266,860 FILING DATE: 2001-02-05
 APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
 APPLICATION NUMBER: GB 24263.6
FILING DATE: 2000-10-04
APPLICATION NUMBER: US 60/236,359
FILING DATE: 2000-09-27
 APPLICATION NUMBER: US 60/207,456
 APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
 FILING DATE:
 FILING DATE:
 FILING DATE: 2001-01-30
 FILING DATE:
 FILING DATE: 2001-01-30 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00666
 APPLICATION NUMBER: PCT/US01/00662
 1357 GCGCCTGGAAGAG 1369
 INFORMATION
 030, Application US/09866108
US20020048800A1
 GCGCCTGGAAGAG 13
 RANK, David R. CHEN, Wensheng
 PENN, Sharron G. HANZEL, David K.
 Yonggang
 2001-01-30
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 0.9%; Score 13;
100.0%; Pred. No.
 2001-05-2
0.9%; Score 13;
100.0%; Pred. No.
 <u>,</u>
 Mismatches
 DB 1; Length 17;
 1.6e+02;
 DB 1;
 ä
 Length 17
 HUMAN HEART
 Indels
 AND MUSCLE
 0
 Gaps
 0
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; ORGANISM: Homo sapiens US-09-866-108-9031
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 RESULT 140
US-09-866-108-9031

 Matches

 PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR PRIOR DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
 Query Match
Best Local Similarity
Matches 13; Conserv
 SOFTWARE: Aec
SEQ ID NO 9031
 Sequence 9031, Application US/09866108 Patent No. US20020048800A1
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 NUMBER OF SEQ ID NOS: 15752
 PRIOR APPLICATION NUMBER: US 60/207,456 PRIOR FILING DATE: 2000-05-26
 FILE REFERENCE
 APPLICANT: GU, Yizhong
 TYPE: DNA
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 PPLICANT: SHANNON, Mark
TILE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 LENGTH:
 FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00668
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00662
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00670
 2397 CGTGGAGGAACTT 2409
 2397 CGTGGAGGAACTT 2409
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 Aeomica Sequence Listing Engine
CGTGGAGGAACTT 16
 G DATE: 2001-01-30
CATION NUMBER: PCT/US01/00663
 CGTGGAGGAACTT 17
 PENN, Sharron HANZEL, David
 CHEN, Wensheng
 RANK, David R
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 AEOMICA-7
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 Score 13; DB 1;
Pred. No. 1.6e+02
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 Length 17;
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 Gaps
 Gaps
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RESULT 141 US-09-866-108-9032

Sequence 9032, Application US/09866108

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 US-09-866-108-9033
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Matches
 Sequence 9033, Application US/09866108 Patent No. US20020048800A1
 SEQ ID NO 9032
 GENERAL INFORMATION:
 Query Match
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggar
APPLICANT: PENN, Sharı
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
 RPPLICANT: SHANNON, MAIK
FITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 PRIOR FILING DATE:
 TYPE: DNA ORGANISM: Homo sapiens
 ENGTH: 17
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 APPLICATION NUMBER: PCT/US01/00666
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: US 60/266,860
 APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 2397 CGTGGAGGAACTT 2409
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00661
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00662
 INFORMATI
 ILING DATE:
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 l Similarity
13; Conserv
INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 US20020048800A3
 GU, Yizhong
 PENN, Sharron G.
HANZEL, David K.
RANK, David R.
CHEN, Wensheng
 RANK, David R. CHEN, Wensheng
 PENN, Sharron G. HANZEL, David K.
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 Yonggang
 Yonggang
 AEOMICA-
 2001-02-05
 2000-09-21
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 2001-01-30
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100.0%; Fi
 0.9%; Score 13; DB 1;
100.0%; Pred. No. 1.6e+0
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 Mismatches
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hes 0;
 Length 17;
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 Gaps
 0
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 RESULT 143
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 ORGANISM: Homo sapiens US-09-866-108-9033
 Sequence 9034, Application US/09866108
Patent No. US20020048800A1
 Matches
 Query Match
Best Local
 SEQ ID NO 9033
 GENERAL
 LENGTH: 17
TYPE: DNA
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 PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga
 PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
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 FILE REFERENCE: AEOMICA-7
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
 PRIOR APPLICATION NUMBER: GB 24263.6 PRIOR FILING DATE: 2000-10-04
 PRIOR APPLICATION NUMBER: US 60/207,456 PRIOR FILING DATE: 2000-05-26
 PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 APPLICANT: SHANNON, Mark
FITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN
 APPLICANT:
 APPLICANT:
 APPLICANT:
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: US 60/236,359
 APPLICATION NUMBER: PCT/US01/00668
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00665
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 APPLICATION NUMBER: US 60/266,860
 FILING DATE: 2000-09-21
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00662
 2397 CGTGGAGGAACTT 2409
 FILING DATE: 2001-01-30
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Similarity 100.0%; Pred. No. 1.6e+
 Aeomica Sequence Listing Engine
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 CGTGGAGGAACTT 14
 PENN, Sharron G. HANZEL, David K.
 RANK, David R. CHEN, Wensheng
 JI, Yonggang
 Conservative
 2001-01-30
PCT/US01/00666
 60/234,687
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 Mismatches
 1.6e+02;
 Length 17;
 HEART AND
 ٥,
 MUSCLE
 Gaps
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RESULT 144
US-09-961-077-748
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 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-9034
 Sequence 748, Application US/09961077 Publication No. US20030014775A1 GENERAL INFORMATION:
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Best Local Similarity
 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 9034
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 NUMBER OF SEQ ID NOS: 15752
 ENGTH: 17
 APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
 CORRESPONDENCE ADDRESS:
 2397 CGTGGAGGAACTT 2409
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
 COMPUTER READABLE FORM:
 NUMBER OF SEQUENCES: 1263
 TITLE
 APPLICANT: Zwick, Michael G.
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00667
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
 FILING DATE:
 13;
 LING DATE: 2001-01-30
 LING DATE: 2001-01-30
 ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
 MEDIUM TYPE: 3.5" Diskette,
 COUNTRY: U.S.A. ZIP: 90071-2066
 CITY: Los Angeles
STATE: California
 QF
G
 0.9%; Score 13; DB 1; larity 100.0%; Pred. No. 1.6e+Conservative 0; Mismatches
 INVENTION: COMPOSITION AND METHODS MODULATION OF GENE EXPRI
 Folkerts, Otto
Merlo, Donald J.
 Skokut, Thomas A. Young, Scott A.
 Merlo,
 Edington, Brent E.
McSwiggen, James A.
 Suite 4700
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 Lining
 Patricia Ann Owens
 IN PLANTS
 1.44 Mb
 ND METHODS FOR GENE EXPRESSION
 1.6e+02;
 Length 17;
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 Gaps
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 RESULT 145
US-09-780-533A-1806/c
 US-09-780-533A-1806
 SOFTWARE: PatentIn version 3.0
SEQ ID NO 1806
LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 Best Local Similarity Matches 13; Conserv
 Sequence 1806, Application US/09780533A Publication No. US20030060611A1
 Query Match
 Query Match 0.9%; Score 13; DB 1; Best Local Similarity 76.9%; Pred. No. 1.6e+02;
 GENERAL INFORMATION:
 Matches
 TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene FILE REFERENCE: MSHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR PILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
 APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: Blatt, Larry APPLICANT: McSwiggen, Jim
 TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 748:
SEQUENCE CHARACTERISTICS:
 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/679,645
FILING DATE: July 12, 1996
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
 ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
 1573 TCCAGCTCCTCCA 1585
 1866 GGTGTCAGAGATG 1878
 SEQUENCE DESCRIPTION: SEQ ID NO: 748:
 REFERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
14
 McSwiggen, Jim
Chowrira, Bharat
Haeberli, Pete
 GGUGUCAGAGAUG 16
 LENGTH: 17 base pairs TYPE: nucleic acid
 CLASSIFICATION: <Unknown>
 STRANDEDNESS: single TOPOLOGY: linear
 APPLICATION NUMBER: US/09/961,077
 TELEFAX: (213) 955-0440
 FILING DATE: 21-Sep-2001
 Conservative
 Conservative
 100.0%;
 0.9%; Score 13; DB 1; L
100.0%; Pred. No. 1.6e+02;
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 Mismatches
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 Length 17;
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RESULT 146 US-09-848-754A-2903

Sequence 2903, Application US/09848754A Publication No. US20030073207A1

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US-09-848-754A-3487
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 RESULT 148
 US-09-848-754A-3486
 RESULT 147
 US-09-848-754A-2903
 US-09-848-754A-3487
 US-09-848-754A-3486
 CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
SOFTWARE: Patentin version 3.0
SEQ ID NO 3487
 Sequence 3487, Application US/09848754A
Publication No. US20030073207A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-I (400/018)
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBHBO0-958-I (400/018)
CURRENT APPLICATION NUMBER: US/99/848,754A
CURRENT FILING DATE: 2001-05-03
 Sequence 3486, Application US/09848754A Publication No. US20030073207A1
 Matches
 Query Match
Best Local Similarity
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 2903
 GENERAL INFORMATION:
 Query Match
Best Local
 Query Match
 Best Local Similarity
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-I (400/018)
FURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 / Match 0.9%;
Local Similarity 76.9%;
nes 10; Conservative
 2273 TCAAGTGGATGGC 2285
 2273 TCAAGTGGATGGC 2285
 10;
 3 UCAAGUGGAUGGC 15
 1 UCAAGUGGAUGGC 13
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 Score 13; DB 1; Lo
Pred. No. 1.6e+02;
3; Mismatches 0;
 Score 13; DB 1;
Pred. No. 1.6e+02
 Score 13;
Pred. No.
 Mismatches
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 DB 1;
 DB 1; Length 17;
 Length 17
 0; Indels
 Length 17;
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 RESULT 149
US-09-776-474-1033
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 US-09-740-332-831
 RESULT 150
US-09-740-332-831/c
 ; OTHER INFORMATION: Description of Artificial Sequence: US-09-776-474-1033
 Sequence 831, Application US/09740332 Publication No. US20030125270A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 1033
 GENERAL INFORMATION:
 Sequence 1033, Application US/09776474
 Matches
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 Best Local
 Query Match
 SEQ ID NO 831
 Best Local Similarity Matches 12; Conserv
 Query Match
 Publication No.
 CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: PatentIn version 3.0
 APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
 NUMBER OF SEQ ID NOS: 2992
 CURRENT APPLICATION NUMBER: US/09/776,474
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR FILING DATE: 2000-03-02
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent
TITLE OF INVENTION: Enzyme
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICANT: Ribozyme
 FILE REFERENCE: MBHB00-955-A (400/008)
 FEATURE:
NAME/KEY: misc_feature
 OTHER INFORMATION: oligonucleotide substrate
 ORGANISM: artificial sequence
 LENGTH: 17
TYPE: RNA
 FEATURE
 TYPE: RNA ORGANISM: Artificial Sequence
 LOCATION:
 ENGTH: 17
1545 GCGGAGACAGGTA 1557
 1730 CCCTGGGAGAAGG 1742
 2273 TCAAGTGGATGGC 2285
 10;
 u
 Similarity
 UCAAGUGGAUGGC 14
 CCCUGGGAGAAGG 17
 Booher, Robert
Holman, Patricia
 Jarvis,
 Fattaey, Ali
 Conservative
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 US20030087847A1
 Pharmaceuticals, Inc
 0.9%;
 0.9%; Score 13;
100.0%; Pred. No.
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 Score 13; DB
Pred. No. 1.6e
1; Mismatches
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 DB 1; L., 1.6e+02;
 DB 1;
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 Length 17;
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 Conditions
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RESULT 153
US-09-817-879-3724
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 RESULT 152
US-09-817-879-831/c
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 US-09-817-879-831
 ; OTHER INFORMATION: oligonucleotide substrate US-09-740-332-3724
 RESULT 151
US-09-740-332-3724
 Sequence 3724, Application US/09817879 Publication No. US20030171311A1 GENERAL INFORMATION:
 GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases

TITLE OF INVENTION: Hepatitis C Virus Infection

FILE REFERENCE: MBHB00-801-F

CURRENT APPLICATION NUMBER: US/09/817,879

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9703

SOFTWARE: Patentin version 3.0

SEQ ID NO 831
 Sequence 3724, Application US/09740332
Publication No. US20030125270A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT PILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: Patentin version 3.0
LENGTH. 3724
 Query Match
Best Local Similarity
 Matches
 Sequence 831, Application US/09817879 Publication No. US20030171311A1
 Best Local Similarity 92.3
Matches 12; Conservative
 Query Match
 LENGTH: 17
TYPE: RNA
ORGANISM: artificial sequence
 LOCATION:
OTHER INFORMATION: oligonucleotide substrate
 NAME/KEY: misc_feature
 TYPE: RNA ORGANISM: artificial sequence
 FEATURE:
 NAME/KEY: misc_feature LOCATION:
 FEATURE:
 ENGTH: 17
 1545 GCGGAGACAGGTA 1557
 1545 GCGGAGACAGGTA 1557
 16
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 GCGGAGACAGGTA 4
 GCGGAGACAGGUÁ 15
 GCGGAGACAGGTA
 Conservative
Pharmaceuticals Inc
 0.9%;
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 Score 13; DB 1; L; Pred. No. 1.6e+02; 0; Mismatches 0;
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 Score 13;
Pred. No.
 Mismatches
 1.6e+02;
 DB 1;
 0; Indels
 Length 17;
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 <u>,,</u>
 or Conditions Relate
 Gaps
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US-09-829-855-177
 RESULT 155
 뭉
 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-230-006;1403
 RESULT 154
US-10-230-006-1403/c
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 US-09-817-879-3724
 Sequence 177, Application US/09829855 Patent No. US20020065609A1 GENERAL INFORMATION:
 NUMBER OF SEQ ID NOS: 2678
SOFTWARE: Patentin version 3.0
SEQ ID NO 1403
LENGTH: 17
 CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3724
LENGTH: 17
TYPE: RNA
 Sequence 1403, Application US/10230006 Publication No. US20030191077A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
Matches 12; Conserv
 Matches
 Query Match
 CURRENT APPLICATION NUMBER: US/09/829,855
CURRENT FILLING DATE: 2001-04-10
PRIOR APPLICATION NUMBER: US 60/196063
PRIOR FILING DATE: 2000-04-10
PRIOR APPLICATION NUMBER: US 60/196258
PRIOR FILING DATE: 2000-04-11
NUMBER OF SEQ ID NOS: 244
 APPLICANT: Matthew, Ashby N.

TITLE OF INVENTION: Methods for the Survey and Genetic Analysis of Populations
FILE REFERENCE: ASHBY-1
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Fosnaugh, Kathy APPLICANT: McSwiggen, Jim TITLE OF INVENTION: METHOD AND REAGENT FOR FILE REFERENCE: 400/056 (MBHB01-1110)
 CURRENT APPLICATION NUMBER: US/10/230,006
CURRENT FILING DATE: 2002-11-18
PRIOR APPLICATION NUMBER: US 60/315,315
PRIOR FILING DATE: 2001-08-28
 TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate TITLE OF INVENTION: Hepatitis C Virus Infection
 FILE REFERENCE: MBHB00-801-F
 FEATURE: NAME/KEY: misc_feature
 OTHER INFORMATION: oligonucleotide substrate
 ORGANISM: artificial sequence
 OCATION:
 / Match 0.9%; So Local Similarity 100.0%; F
 1573 TCCAGCTCCTCCA 1585
 1545 GCGGAGACAGGTA 1557
 14 TCCAGCTCCTCCA 2
 3 GCGGAGACAGGUA 15
PatentIn version 3.1
 Conservative
 0.9%;
 Score 13; ; Pred. No.
 1; Mismatches
 Score 13; DB 1;
Pred. No. 1.6e+02;
 Mismatches
 DB 1; Le
 THE TREATMENT OF ASTHMA AND ALLERGIC CONDI
 Length 17;
 Length 17
 Indels
 0
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 Gaps
 Gaps
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US-09-866-108-653/c
 밁
 ; OTHER INFORMATION: Uncultured Acidobacterium Sub.Div-1 US-09-829-855-177
 ; ORGANISM: Homo sapiens US-09-866-108-653
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR PPLICATION NUMBER: GB 24263.6
PRIOR PPLICATION NUMBER: GB 24263.6
PRIOR PPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR PILING DATE: 2001-01-30
PRIOR PR
 RESULT 156
 SEQ ID NO 177
LENGTH: 16
 Sequence 653, Application US/09866108 Patent No. US20020048800A1
 Matches
 Query Match
Best Local Similarity
Best Local Similarity
 Query Match
 APPLICANT:
 PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga
APPLICANT: PENN, Shar
 APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART
 NUMBER OF SEQ ID NOS: 15752
 LENGTH: 17
TYPE: DNA
 SOFTWARE: Aeomica Sequence Listing Engine
 ORGANISM: Unknown
 TYPE: DNA
 APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00662
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663
 1511 CGGCTGTGCACAAGCT 1526
 INFORMATI
 14; Conservative
 1 CGGCTGTGCCGAAGCT 16
 JI, Yonggang
PENN, Sharron G.
HANZEL, David K.
 RANK, David R
 0.9%;
0.9%;
 Score 12.8;
Pred. No. 1.
Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 DB 1;
.5e+02;
 Length 17;
 Length 16
 Indels
 0
 AND MUSCLE
 Gaps
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 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/USO1/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00667
PRIOR PILING DATE: 2001-01-30
PRIOR PPLICATION NUMBER: PCT/USO1/00664
PRIOR PPLICATION NUMBER: PCT/USO1/00669
PRIOR PPLICATION NUMBER: PCT/USO1/00669
PRIOR PPLICATION NUMBER: PCT/USO1/00669
PRIOR APPLICATION NUMBER: PCT/USO1/00665
PRIOR PPLICATION NUMBER: PCT/USO1/00666
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR APPLICATION NUMBER: PCT/USO1/00662
PRIOR APPLICATION NUMBER: PCT/USO1/00662
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR PILING DATE: 2001-01-30
PRIOR PRIOR PILING DATE: 2000-09-21
PRIOR PILING DATE: 2000-09-21
 RESULT 157
US-09-866-108-654/c
 맑
 á
 ; ORGANISM: Homo sapiens US-09-866-108-654
 SOFTWARE: Aeor
; SEQ ID NO 654
; SEQIPH: 17
 GENERAL
 Sequence 654, Application US/09866108 Patent No. US20020048800A1
 Matches
 Matches
 Query Match
Best Local Similarity
 APPLICANT: SHANNON, MAYK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT: GU, YI
APPLICANT: JI, Y
APPLICANT: PENN,
 PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2001-02-05
 NUMBER OF SEQ ID NOS: 15752
 TYPE: DNA
 1566 GGCTGAGTCCAGCTCC 1581
 1566 GGCTGAGTCCAGCTCC 1581
 INFORMATION:
 17
16
 14;
 14;
 Aeomica Sequence Listing Engine
GGCTGAGTCCGGGTCC 1
 JI, Yonggang PENN, Sharro
 RANK, David k. CHEN, Wensheng
 HANZEL, David
 Conservative
 Conservative
 Sharron
 2001-02-05
 0.9%;
 60/266,860
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 ;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 Mismatches
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 Length 17;
 Indels
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 Gaps
 Gaps
 0
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RESULT 158 US-09-866-108-1107 ; Sequence 1107, Application US/09866108

FILE REFERENCE: AEOMICA-7

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US-09-866-108-1108
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 ; LENGTH: 17
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-1107
 Sequence 1108, Application US/09866108 Patent No. US20020048800A1
 SOFTWARE: Aeomica Sequence SEQ ID NO 1107
 Matches
 Query Match
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 APPLICANT:
 PRIOR APPLICATION NUMBER: US 60/266,860 PRIOR FILING DATE: 2001-02-05
 APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
 APPLICANT:
 NUMBER OF SEQ ID NOS: 15752
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 PRIOR
 PRIOR APPLICATION NUMBER: GB 24263.6 PRIOR FILING DATE: 2000-10-04
 PRIOR APPLICATION NUMBER: US 60/207,456 PRIOR FILING DATE: 2000-05-26
 APPLICATION NUMBER: PCT/US01/00667
 APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
 APPLICATION NUMBER: US 60/236,359 FILING DATE: 2000-09-27
 FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
 APPLICATION NUMBER: PCT/US01/00669
 FILING DATE: 2001-01-30 APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00663 FILING DATE: 2001-01-30
 1725 CAAGCCCCTGGGAGAA 1740
 APPLICATION NUMBER: PCT/US01/00670
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00665
 FILING DATE:
 FILING DATE:
 INFORMATION:
 PPLICATION NUMBER: PCT/US01/00668
TLING DATE: 2001-01-30
 INFORMATION
 LING DATE: 2001-01-30
 LING DATE:
 N
 14;
TIVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 US20020048800A1
 GU, Yizhong
 JI, Yonggang
PENN, Sharron G.
HANZEL, David K.
RANK, David R.
CHEN, Wensheng
 CAAATCCCTGGGAGAA 17
 HANZEL,
 PENN
 Conservative
 Yizhong
 Yonggang
 , Sharron G.
EL, David K.
 2001-01-30
 2001-01-30
 2001-01-30
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 2001-05-25
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 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 Length 17,
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; LENGTH: 17
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-1108
 US-09-866-108-1112
 SOFTWARE: Aeomica Sequence
SEQ ID NO 1108
 GENERAL
 Sequence 1112, Application US/09866108 Patent No. US20020048800A1
 Matches
 Query Match
 APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: ABOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2000-05-26
 APPLICANT:
 PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
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 APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharro
 PRIOR FILING DATE: 2001-02-
NUMBER OF SEQ ID NOS: 15752
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 PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
 PRIOR FILING DATE: 2001-01 US01/00662
 PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: US 60/236,359 PRIOR FILING DATE: 2000-09-27
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT:
 PRIOR APPLICATION NUMBER: PCT/US01/00663
 PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00666
 Local Similarity les 14; Conserv
 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00666
 APPLICATION NUMBER: US 60/266,860 FILING DATE: 2001-02-05
 APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
 APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668
 1725 CAAGCCCCTGGGAGAA 1740
 INFORMATION:
 1 CAAATCCCTGGGAGAA 16
 PENN, Sharron G.
HANZEL, David K.
RANK, David R.
CHEN, Wensheng
 Conservative
 2000-09-
 0.9%;
 Listing Engine
 °.
 Score 12.8;
Pred. No. 1
 Mismatches
 1.7e+02;
 DB 1;
 Z
 HUMAN
 Length 17;
 Indels
 HEART AND MUSCLE
 0,
 Gaps
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 ; ORGANISM: Homo sapiens
US-09-866-108-1112
 SEQ ID NO 1112
 Query Match
Best Local Similarity
 Sequence 1113, Application US/09866108 Patent No. US20020048800A1
 Matches
 APPLICANT:
APPLICANT:
APPLICANT:
 NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engine
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 LENGTH: 17
TYPE: DNA
 PRIOR APPLICATION NUMBER: US 60/266,860 PRIOR FILING DATE: 2001-02-05
 FILE REFERENCE: AEOMICA-7
 APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
 APPLICANT:
 APPLICANT: GU, Yizhong
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
 APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
 APPLICATION NUMBER: PCT/US01/00663 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00667 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00666 FILING DATE: 2001-01-30
 1730 CCCTGGGAGAAGGTTG 1745
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00661
 FILING DATE: 2001-01-30
 APPLICATION NUMBER:
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00665
 INFORMATION:
 14; Conservative
 2 CCCTGGGAGAAAGTGG 17
 RANK, David R. CHEN, Wensheng
 PENN, Sharron G. HANZEL, David K.
 Yonggang
NUMBER: PCT/US01/00668
 2001-01-30
 0.9%;
 2001-05-25
 PCT/US01/00662
 Score 12.8; DB
Pred. No. 1.7e+C
0; Mismatches
 1.7e+02
 Ä
 HUMAN HEART
 Length 17;
 AND MUSCLE
 0,
 Gaps
 0,
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 á
 US-09-866-108-1175
 ; ORGANISM: Homo sapiens US-09-866-108-1113
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PELING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PRILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00661
 SEQ ID NO 1113
 Sequence 1175, Application US/09866108 Patent No. US20020048800A1
 Matches
 Query Match
 APPLICANT: GU,
APPLICANT: JI,
APPLICANT: PEN
APPLICANT: HAN
APPLICANT: RAN
 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 SOFTWARE: Aeomica Sequence Listing Engine
 PRIOR FILING DATE: 2000-09-21
PRIOR PELICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
 PRIOR APPLICATION NUMBER: PCT/US01/00670
 APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 PRIOR
 LENGTH:
 Local Similarity
 APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: US 60/234,687
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
 FILING DATE:
 1730 CCCTGGGAGAAGGTTG 1745
 INFORMATION:
 14;
 CCCTGGGAGAAAGTGG
 RANK, David R. CHEN, Wensheng
 PENN, Sharron G.
HANZEL, David K.
 JI, Yonggang
 Conservative
 Yizhong
 2001-01-30
 2001-01-30
 2001-01-30
 0.9%;
-01 - 30
 16
 <u>.</u>
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 2
 DB 1;
 Length 17;
 Indels
 0,
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 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 1175
LENGTH: 17
 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 1176
 Sequence 1176, Application Patent No. US20020048800A1
 Query Match
TYPE: DNA
ORGANISM: Homo sapiens
-09-866-108-1176
 Matches
 LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
-09-866-108-1175
 PRIOR FILING DATE: 2000-02-2.
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
NUMBER OF SEQ ID NOS: 15752
 ENERAL INFORMATION:
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
 APPLICANT:
 APPLICANT:
 PPLICANT:
 Match 0.9%;
Local Similarity 87.5%;
 R OF SEQ ID NOS: 15752
 APPLICATION NUMBER: GB 24263.6
FILING DATE: 2000-10-04
APPLICATION NUMBER: US 60/236,359
 APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30 APPLICATION NUMBER: US
 APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00667
 FILING DATE:
 FILING DATE: 2000-09-21
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00664 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00663
 2425 GAAGGACACAGAATGG 2440
 LING DATE: 2001-01-30
 LING DATE: 2001-01-30
 LING DATE: 2001-01-30
 LING DATE: 2001-01-30
 N
 GAAGGACAAAGAAGGG 17
 HANZEL, David K.
 JI, Yonggang
 Conservative
 AEOMICA-7
 2001-01-30
 2001-01-30
 2000-09-
 US 60/234,687
 US/09866108
 0
 Score 12.8;
Pred. No. 1.
 Mismatches
 .7e+02
 DB 1;
 IN HUMAN HEART AND MUSCLE
 Length 17;
 Indels
 0
 0
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 ; ORGANISM: Homo sapiens US-09-866-108-1571
 RESULT 164 .
US-09-866-108-1571
 맑
 SEQ ID NO 1571
 Query Match
Best Local Similarity
Matches 14; Conserv
 Query Match
 Matches
 Sequence 1571, Application US/09866108 Patent No. US20020048800A1
 LENGTH: 17
TYPE: DNA
 PRIOR
PRIOR
 APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
 PRIOR FILING DATE: 200 NUMBER OF SEQ ID NOS:
 PRIOR
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 PRIOR APPLICATION NUMBER: US 60/266,860
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 PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 2001-0:
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00666
 RIOR APPLICATION NUMBER: PCT/US01/00670
 RIOR APPLICATION NUMBER: PCT/US01/00661
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 Local Similarity
 FILING DATE: 2001-01-30
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00662
 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00669
 1503 CAGCCAGCCGGCTGTG 1518
 FILING DATE: 2000-09-21
 APPLICATION NUMBER: US
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00665
 FILING DATE:
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00664
 2425 GAAGGACACAGAATGG 2440
 INFORMATION:
 14;
 Aeomica Sequence Listing Engine
CAGCCAGCCTGCTCTG 17
 PENN, Sharron G
HANZEL, David K
RANK, David R.
CHEN, Wensheng
 GAAGGACAAAGAAGGG 16
 Conservative
 Conservative
 Yonggang
 2001-02-05
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 0.9%;
87.5%;
 0.9%;
 PCT/US01/00667
 60/234,687
 0;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8;
 Mismatches
 Mismatches
 No.
 1.7e+02;
 DB 1;
 Ä
 1,
 HUMAN HEART
 Length 17;
 Length 17;
 Indels
 Indels
 0
 AND MUSCLE
 0
 Gaps
 Gaps
 0,
 0
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RESULT 165

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; SEQ ID NO 1572
; LENGTH: 17
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-1572
 В
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 US-09-866-108-1572
 US-09-866-108-6176
 RESULT 166
 Query Match
Best Local S
Matches 14
 GENERAL INFORMATION:
 Sequence 6176,
 Sequence 1572, Application US/09866108 Patent No. US20020048800A1
 Patent No.
 APPLICANT: GU, Yizhong APPLICANT: JI, Yonggar
 PRIOR FILING DATE: 2000-10 PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 2000-05
 PRIOR APPLICATION NUMBER: US 60/207,456 PRIOR FILING DATE: 2000-05-26 PRIOR APPLICATION NUMBER: GB 24263.6
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT: SHANNON, MARK
IITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 APPLICANT:
 APPLICANT: GU, Yizhong
 SOFTWARE: Aeomica Sequence Listing Engine
 Local Similarity
nes 14; Conserv
 APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
 FILING DATE:
 APPLICATION NUMBER: US 60/266,860
 APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00664 FILING DATE: 2001-01-30
 FILING DATE:
 FILING DATE: 2000-09-21
 APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00665
 1503 CAGCCAGCCGGCTGTG 1518
 APPLICATION NUMBER: US 60/234,687
 FILING DATE:
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663
 INFORMATION:
 LING DATE:
 LING DATE:
 LING DATE:
 ...
 176, Application US/09866108
US20020048800A1
 CAGCCAGCCTGCTCTG 16
JI, Yonggang
PENN, Sharron G.
HANZEL, David K.
RANK, David R.
 CATION NUMBER: PCT/US01/00661
 CHEN, Wensheng
 HANZEL,
 PENN
 Conservative
 , Yonggang
NN, Sharron G.
NZEL, David K.
 AEOMICA-7
 2001-01-30
 2000-10-04
NUMBER: US 60/236,359
2000-09-27
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 2001-02-05
 2001-01-30
 2001-01-30
 2001-01-30
 0.9%;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 2;
 Length 17;
 Indels
 0
 Gaps
 0
```

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밁
 S
 US-09-866-108-6177
 RESULT 167
 ; ORGANISM: Homo sapiens US-09-866-108-6176
 SOFTWARE: Aeon
SEQ ID NO 6176
 Sequence 6177, Application US/09866108 Patent No. US20020048800A1
 Query Match
Best Local Similarity
 GENERAL
 Matches
 APPLICANT:
APPLICANT:
APPLICANT:
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT: SHANNON, MAXK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
 APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 APPLICANT:
 PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
 PRIOR APPLICATION NUMBER: GB 24263.6 PRIOR FILING DATE: 2000-10-04
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT:
 APPLICANT: GU, Yizhong
 NUMBER OF SEQ ID NOS: 15752
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: US 60/266,860
 PRIOR APPLICATION NUMBER: PCT/US01/00668 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00667
 APPLICANT:
 TYPE: DNA
 ENGTH:
 APPLICATION NUMBER: PCT/US01/00661
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00663
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2000-10-04
 APPLICATION NUMBER: US 60/234,687
 FILING DATE: 2001-01-30
 1784 ACAAAGACAAGCCCAA 1799
 INFORMATION:
 14;
 N
 Aeomica Sequence Listing Engine
 HANZEL, David K
RANK, David R.
CHEN, Wensheng
 ACAGAGCCAAGCCCAA 17
 CATION NUMBER: PCT/US01/00670
 CATION NUMBER: PCT/US01/00665
 JI, Yonggang
PENN, Sharron
 CHEN,
 Conservative
 Wensheng
 2001-01-30
 2000-09-21
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 2001-02-05
 0.9%;
 ჯ ი
 0
 Score 12.8;
Pred. No. 1
 Mismatches
 1.7e+02
 DB 1;
 Length 17,
 Indels
 <u>,</u>
 Gaps
 0,
```

NUMBER: US 60/236,359

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RESULT 168
US-09-866-108-6311
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 ; SEQ ID NO 6177
; LENGTH: 17
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-6177
 Sequence 6311, Application US/09866108 Patent No. US20020048800A1
 Matches
 Query Match
Best Local Similarity 87.5%;
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR PRILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
 APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
FILE REFERENCE: AEOMICA-7
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT:
 PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggan
 SOFTWARE: Aeomica Sequence Listing Engine
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00661
FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
 1784 ACAAAGACAAGCCCAA 1799
 APPLICATION NUMBER: US 60/266,860
 FILING DATE: 2000-09-21
 APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00664 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00666 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00667
 INFORMATION
 14;
 LING DATE: 2001-01-30
 LING DATE: 2001-01-30
 PENN, Sharron G.
HANZEL, David K.
RANK, David R.
CHEN, Wensheng
 ACAGAGCCAAGCCCAA 16
 Conservative
 Yonggang
 2000-09-
 2001-05-25
 Score 12.8; D
Pred. No. 1.7e
0; Mismatches
 .7e+02;
 DB 1;
 HUMAN HEART
 Length 17;
 Indels
 AND MUSCLE
 0
 Gaps
 0
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 US-09-866-108-6311
 US-09-866-108-6313
 RESULT 169
 PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/USO1/00666
PRIOR TILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00667
PRIOR APPLICATION NUMBER: PCT/USO1/00667
PRIOR FILING DATE: 2001-01-30
PRIOR PRILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR APPLICATION NUMBER: PCT/USO1/00665
 Best Loc
Matches
 SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 6311
LENGTH: 17
 Sequence
 Query Match
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00662
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00661
 APPLICANT:
 APPLICANT:
APPLICANT:
 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
 CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 APPLICANT: GU
APPLICANT: J:
 NUMBER OF SEQ ID NOS: 15752
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
 PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
 APPLICANT: SHANNON, Mark
FITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 TYPE: DNA
ORGANISM: Homo sapiens
 PRIOR APPLICATION NUMBER: PCT/US01/00670
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00662
 PRIOR FILING DATE: 2001-01-30
 PRIOR
 Local Similarity
les 14; Conserv
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30 APPLICATION NUMBER: PCT/US01/00663
 ŏ.
 2090 GCACCTACCAGCTGGC 2105
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00665
 INFORMATION
 6313,
 N
 313, Application US/09866108
US20020048800A1
 GCACCTCCCAGCAGGC 17
 PENN, Sharron
HANZEL, David
 CHEN, Wensheng
 RANK, David R.
 Yizhong
 Conservative
 Yonggang
 0.9%; Score 12.8;
87.5%; Pred. No. 1.
 <u>ج</u> ۵
 Mismatches
 DB 1;
 Length 17;
 0
 Gaps
```

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밁
 8
 ; ORGANISM: Homo sapiens US-09-866-108-6313
 US-09-866-108-6531
 Sequence 6531, Application US/09866108 Patent No. US20020048800A1
 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 6313
 GENERAL INFORMATION:
 Query Match
Best Local
 Matches
 APPLICANT:
 PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
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 NUMBER OF SEQ ID NOS: 15752
 PRIOR APPLICATION NUMBER: US 60/266,860 PRIOR FILING DATE: 2001-02-05
 PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2000-09-21
 PRIOR
 PRIOR
 PRIOR APPLICATION NUMBER: PCT/US01/00666
 PRIOR APPLICATION NUMBER: GB 24263.6 PRIOR FILING DATE: 2000-10-04
 PRIOR FILING DATE:
 CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 APPLICANT: GU, Yizhong
 TYPE: DNA
 PPLICANT:
 ENGTH: 17
 APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: US 60/236,359 FILING DATE: 2000-09-27
 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
 2091 CACCTACCAGCTGGCC 2106
 APPLICATION NUMBER: PCT/US01/00663
 14;
 ب
 Similarity
Aeomica
 CACCICCCAGCAGGCC 16
 HANZEL, David K.
 PENN,
 Conservative
 Yonggang
 AEOMICA-7
Sequence
 2000-05-
 0.9%;
 PCT/US01/00670
Listing Engine
 60/234,687
 0,
 Score 12.8;
Pred. No. 1.
 Mismatches
 DB 1;
 Length 17;
 Indels
 0,
 Gaps
 0;
```

```
; LENGTH: 17
; TYPE: DNA
; ORGANISM: Homo s
US-09-866-108-6532
 δ
 밁
 Ş
 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-6531
 US-09-866-108-6532
 RESULT 171
 PRIOR APPLICATION NUMBER: PCT/USO1/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR APPLICATION NUMBER: PCT/USO1/00662
PRIOR APPLICATION NUMBER: PCT/USO1/00662
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00661
 PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
 Query Match
Best Local S
Matches 14
 Query Match
Best Local S
Matches 14
 SEQ ID NO 6532
LENGTH: 17
 Sequence 6532, Application US/09866108 Patent No. US20020048800A1
 APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANKON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence
 PRIOR APPLICATION NUMBER: US 60/207,456 PRIOR FILING DATE: 2000-05-26
 PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
 APPLICANT:
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: PCT/US01/00670
 PRIOR FILING DATE: 2001-01-30
 0.9%;
Local Similarity 87.5%;
les 14; Conservative
1359 GCCTGGAAGAGAAAAG 1374
 FILING DATE:
 1359 GCCTGGAAGAAAAAG 1374
 INFORMATION:
 2 GCCTGGTAGAGACAAG 17
 Similarity
 Aeomica Sequence Listing Engine
 CATION NUMBER: US 60/234,687
 JI, Yonggang
PENN, Sharron G.
HANZEL, David K.
 Yizhong
 Conservative
 sapiens
 2001-01-30
 2001-02-05
 0.9%;
 Score 12.8;
Pred. No. 1.
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 1.7e+02;
ches 2;
 DB 1;
 Length 17;
 Length 17;
 Indels
 0
 0,
 Gaps
 MUSCLE
 0
 0
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GCCTGGTAGAGACAAG 16

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 á
 US-09-866-108-7346
 RESULT 173
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; ORGANISM: Homo sapiens
US-09-866-108-7345
 US-09-866-108-7345
 Query Match
Best Local Similarity 87.5
14; Conservative
 GENERAL INFORMATION:
 Sequence 7346, Application US/09866108 Patent No. US20020048800A1
 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 7345
 Sequence 7345, Application US/09866108
 Patent No. US20020048800A1
 PRIOR
 PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT: GU, Yizhong APPLICANT: JI, Yongga
 APPLICANT:
 NUMBER OF SEQ ID NOS: 15752
 FILE REFERENCE:
 APPLICANT:
 APPLICANT:
 APPLICANT:
 PLICANT: SHANNON, Mark
[TLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED]
 ENGTH:
 APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
 FILING DATE: 2000-10-04
APPLICATION NUMBER: US 60/236,359
FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00661
FILING DATE: 2001-01-30
 2524 AAGCAGTTGGTAGAAG 2539
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00665
 INFORMATION:
 LING DATE: 2001-01-30
 LING DATE: 2001-01-30
 2 AAACAGTTGGAAGAAG 17
GU, Yizhong
JI, Yonggang
 RANK, David R. CHEN, Wensheng
 HANZEL,
 Yonggang
 AEOMICA-7
 Sharron
 0.9%;
 2001-05-25
 ჯ ი
 0;
 Score 12.8;
Pred. No. 1
 Mismatches
 .7e+02;
 DB 1;
 IN HUMAN HEART AND MUSCLE
 Length 17;
 0;
 Gaps
 0
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RESULT 174
US-09-866-108-7672/c
 밁
 Ś
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 PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR APPLICATION NUMBER: PCT/US01/00661
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PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
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PRIOR PILING DATE: 2001-01-30
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PRIOR FILING DATE: 2000-05-26
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PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
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PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
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PRIOR PILING DATE: 2001-01-30
PRIOR PRIOR PILING DATE: 2001-01-30
 ; SOFTWARE: Aeom
; SEQ ID NO 7346
; LENGTH: 17
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
 Query Match
Best Local Similarity
Matches 14; Conserv
 Sequence 7672, Application US/09866108 Patent No. US20020048800A1
 GENERAL INFORMATION:
 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
 APPLICANT:
 NUMBER OF SEQ ID NOS: 15752
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
 PRIOR FILING DATE: 2001-02-05
 FILE REFERENCE:
 TYPE: DNA
 FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
 APPLICATION NUMBER: US 60/234,687
 2524 AAGCAGTTGGTAGAAG 2539
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 I: SHANNON, Mark
INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 Aeomica Sequence Listing Engine
 AAACAGTTGGAAGAAG 16
 HANZEL, David K
RANK, David R.
CHEN, Wensheng
 0.9%;
ilarity 87.5%;
Conservative
 AEOMICA-7
 Score 12.8;
Pred. No. 1.
 Mismatches
 DB 1;
.7e+02;
 7
 IN HUMAN HEART AND MUSCLE
 Length 17;
 Indels
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 Gaps
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 US-09-866-108-7674/c
 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-7672
 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 7672
 Query Match
Best Local
 Matches
 Patent No.
 FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggar
APPLICANT: PENN, Sharı
APPLICANT: HANZEL, Dav
 PRIOR APPLICATION NUMBER: US 60/207,456 PRIOR FILING DATE: 2000-05-26
 APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MAIK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 NUMBER OF SEQ ID NOS: 15752
 ENGTH:
 APPLICATION NUMBER: PCT/US01/00666
 2659 TCTGTTTTTTCTCCAG 2674
 FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
 APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00667
 FILING DATE: 2000-05-26
APPLICATION NUMBER: GB 24263.6
FILING DATE: 2000-10-04
APPLICATION NUMBER: PCT/US01/00664
 FILING DATE:
 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00661
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00663
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: PCT/US01/00666
 APPLICATION NUMBER: US 60/236,359
 APPLICATION NUMBER:
 FILING DATE: 2001-01-30
 FILING DATE:
 FILING DATE:
 FILING DATE:
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 7674, Application US/09866108

b. US20020048800A1
 17 TÉTÉCTTCTTCTCAG 2
 LING DATE: 2001-01-30
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 Similarity
 Conservative
 Yonggang
 , Sharron G.
EL, David K.
 2001-01-30
 2000-09-27
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
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 PCT/US01/00667
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 Score 12.8; DB 1; Length 17; Pred. No. 1.7e+02; o; Mismatches 2; Indels
 0
 0
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 ફ
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; ORGANISM: Homo sapiens
US-09-866-108-7674
 PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR PPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR PPLICATION NUMBER: PCT/US01/00664
PRIOR PPLICATION NUMBER: PCT/US01/00669
PRIOR PPLICATION NUMBER: PCT/US01/00669
PRIOR PPLICATION NUMBER: PCT/US01/00665
PRIOR PPLICATION NUMBER: PCT/US01/00665
PRIOR PPLICATION NUMBER: PCT/US01/00668
PRIOR PPLICATION NUMBER: PCT/US01/00668
PRIOR PPLICATION NUMBER: PCT/US01/00668
PRIOR PRIOR DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
 SEQ ID NO 7674
 GENERAL
 Sequence 9074, Application Patent No. US20020048800A1
 Matches
 Query Match
Best Local
 APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2000-05-26
PRIOR PILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
 APPLICANT:
 APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga
APPLICANT: PENN, Shar
 PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
 NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence
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 PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE:
 APPLICANT:
 PRIOR
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 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: US 60/266,860 FILING DATE: 2001-02-05
 APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00662
 APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00665
 FILING DATE:
APPLICATION NUMBER: PCT/US01/00662
 2658 TTCTGTTTTTCTCCA 2673
 FILING DATE: 2001-01-30
 INFORMATION:
 16 TTCTGCTTCTTCTCCA 1
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 Aeomica Sequence Listing Engine
 HANZEL, David R. RANK, David R.
 PENN, Sharron
 CHEN, Wensheng
 Application
 Conservative
 Yonggang
 2001-01-30
 2001-01-30
 0.9%;
 <u>ج</u> 0
 US/09866108
 0,
 Score 12.8;
Pred. No. 1
 Mismatches
 1.7e+02;
 DB 1;
 IN HUMAN HEART AND MUSCLE
 Length 17;
 Indels
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S
 밁
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; ORGANISM: Homo sapiens
US-09-866-108-9074
 US-09-866-108-9076
 Query Match
Best Local Similarity
Matches 14; Conserv
 Sequence 9076, Application US/09866108 Patent No. US20020048800A1
 SOFTWARE: Acomica Sequence Listing Engine SEQ ID NO 9074
 PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
 APPLICANT: SHANNON, MAIK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
 NUMBER OF SEQ ID NOS: 15752
 APPLICANT: GU, Yizhong APPLICANT: JI, Yongga
 APPLICANT:
 ENGTH:
 FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
 FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00666
 FILING DATE: 2000-10-04
 APPLICATION NUMBER: US 60/236,359
 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
 FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
 APPLICATION NUMBER: US 60/234,687
 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
 APPLICATION NUMBER: PCT/US01/00661
 APPLICATION NUMBER: PCT/US01/00662 FILING DATE: 2001-01-30
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00665
 APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
 FILING DATE:
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
 APPLICATION NUMBER: PCT/US01/00667
 FILING DATE:
 APPLICATION NUMBER: GB 24263.6
 1397 ACCTGGAGATAGCCAT 1412
 APPLICATION NUMBER: US 60/234,687
 FILING DATE:
 INFORMATION:
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 CHEN, Wensheng
 RANK, David R
 HANZEL,
ID NOS: 15752
 Conservative
 Yonggang
 Sharron
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
 0.9%;
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 ٥,
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 1:
 IN HUMAN HEART AND MUSCLE
 Length 17;
 Indels
 <u>.</u>.
 Gaps
 <u>.</u>
```

```
APPLICANT: HANZEL, David K.

APPLICANT: RANK, David R.

APPLICANT: CHEN, Wensheng

APPLICANT: SHANNON, MARK

TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE

FILE REFERENCE: AEOMICA-7

CURRENT APPLICATION NUMBER: US/09/866,108

CURRENT FILING DATE: 2001-05-25

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR FILING DATE: 2000-05-26

PRIOR FILING DATE: 2000-10-04

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: WS 60/236,359
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US-09-866-108-9076
 US-09-866-108-10700
 RESULT 178
 US-09-866-108-10700
 NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence
SEQ ID NO 10700
LENGTH: 17
 SOFTWARE: Aeomica
SEQ ID NO 9076
LENGTH: 17
Query Match 0.9%;
Best Local Similarity 87.5%;
Matches 14; Conservative
 Sequence
 Matches
 Query Match
Best Local Similarity
 Sequence 10700, Application US/09866108
Patent No. US20020048800A1
 PRIOR FILING DATE: 2000-09-21 PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2001-02-05
 APPLICANT:
 APPLICANT: GU
 PRIOR APPLICATION NUMBER: PCT/US01/00661
 PRIOR APPLICATION NUMBER: PCT/US01/00662
 PRIOR APPLICATION NUMBER: PCT/US01/00663
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00664 PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: PCT/US01/00665
 ORGANISM: Homo sapiens
 APPLICATION NUMBER: US 60/234,687
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00670
 FILING DATE: 2001-01-30
 1398 CCTGGAGATAGCCATT 1413
 INFORMATION:
 14;
 1 CCTGGAGACATCCATT 16
 PENN,
 , Yizhong
 Conservative
 Yonggang
N, Sharron
 sapiens
 Sequence Listing Engine
 2001-01-30
 0.9%;
 Listing Engine
 60/266,860
 0
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
D; Mismatches
 Score 12.8;
Pred. No. 1
 Mismatches
 DB 1;
.7e+02;
 Length
 Length
 Indels
 17;
 ٥.
 Gaps
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Indels

0

Gaps

1833 AGATGATGCCACAGAG

1848

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RESULT 180
US-09-864-785-664/c
; Sequence -664, Application US/09864785
; Patent No. US20020177568A1
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 엉
 US-09-866-108-10701
 US-09-866-108-10701
 SEQ ID NO 10701
 Best Local
 Query Match
 CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
 SOFTWARE: Aeomica Sequence
 APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
 LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
 APPLICANT: GU, Yizhong
 FILING DATE: 2000-05-26
APPLICATION NUMBER: GB 24263.6
 APPLICATION NUMBER: US 60/236,359 FILING DATE: 2000-09-27
 APPLICATION NUMBER: US 60/266,860
 APPLICATION NUMBER: PCT/US01/00666
 FILING DATE: 2000-10-04
 FILING DATE: 2001-01-30 APPLICATION NUMBER: US 60/234,687
 APPLICATION NUMBER: PCT/US01/00670
 FILING DATE: 2001-01-30
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 APPLICATION NUMBER: PCT/US01/00667
 FILING DATE: 2001-01-30
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 APPLICATION NUMBER: PCT/US01/00668
 APPLICATION NUMBER: PCT/US01/00665
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00669
 FILING DATE:
 APPLICATION NUMBER: PCT/US01/00664
 [LING DATE: 2001-01-30
 LING DATE: 2001-01-30
 LING DATE: 2000-09-21
 LING DATE:
 0701, Application US/09866108
US20020048800A1
 Similarity
 ICATION NUMBER: PCT/US01/00662
 HANZEL,
 Conservative
 Yonggang
 ; Sharron G.
EL, David K.
 2001-02-05
 2001-01-30
 2001-01-30
 2001-01-30
 2001-01-30
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 Score 12.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 2;
 Length 17;
 0;
 Gaps
 0,
```

APPLICANT:

Karpeisky, Alex Adamic, Jasenka Matulic

Sweedler, Dave

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 US-09-825-805-782
 밁
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 US-09-864-785-1617
 RESULT 181
US-09-864-785-1617
 US-09-864-785-664
 RESULT 182
 Sequence 782, Application US/09825805
Publication No. US20030004122A1
GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 1617
 SEQ ID NO 664
LENGTH: 17
 Query Match
 GENERAL INFORMATION:
 Matches
 Query Match
 GENERAL INFORMATION:
 Matches
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 Sequence 1617, Application US/09864785
Patent No. US20020177568A1
 APPLICANT: Draper, Ken
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate-
TITLE OF INVENTION: Levels of NF-Kappa B
FILE REFERENCE: 400/022 (MBHB00-812-D)
CURRENT FILING INVENTION NUMBER: US/09/864,785
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 3929
 APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: Beigelman, Leo
 CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 3929
SOFTWARE: PatentIn version 3.0
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Stinchcomb, Dan
APPLICANT: Draper, Ken
APPLICANT: Draper, Ken
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate-
TITLE OF INVENTION: Levels of NF-Kappa B
 APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Stinchcomb, Dan
 TITLE OF INVENTION: Levels of NF-Kappa B FILE REFERENCE: 400/022 (MBHB00-812-D)
 APPLICANT:
 ORGANISM: Artificial Sequence
 TYPE: RNA
 OTHER INFORMATION: Description of Artificial Sequence:
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 TYPE: RNA
 OTHER INFORMATION: Description of Artificial Sequence:
 FEATURE:
 LENGTH:
 ENGTH:
 POCAT
 Local Similarity
 2005 GCCCGGAGGCCACCCG 2020
 1398 CCTGGAGATAGCCATT 1413
 16 CCTGGAGAGAGCCAGT 1
 14;
 Similarity
 GCCCAGAGGCCCCCCG 16
Beaudry, Amber
 Beigelman,
 Conservative
 Conservative
 0.9%;
 0.9%;
 0
 0
 Score 12.8;
Pred. No. 1.
 Score 12.8; DB 1.
Pred. No. 1.7e+02.
 Mismatches
 1.7e+02
 DB 1;
 DB 1;
 Length 17;
 Length 17;
 Indels
 Nucleic Acid
 Nucleic
 0
 <u>.</u>
 Acid
 Gaps
 Gaps
 0
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 5
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 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-818-875-1759
 Query Match
Best Local S
 PRIOR APPLICATION NUMBER: 09/474,432
PRIOR FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 09/301,511
PRIOR FILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: 09/186,675
PRIOR FILING DATE: 1998-11-04
PRIOR APPLICATION NUMBER: 60/083,727
 SEQ ID NO 1759
 Sequence 1759, Application US/09818875 Publication No. US20030051270A1
 Query Match
Best Local 9
 Matches
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
-09-825-805-782
 PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR FILING DATE: 2000-10-30
 APPLICANT: Gamper, Howard B.
APPLICANT: Rice, Michael B.
TITLE OF INVENTION: Target C. Chromosomal Genomic Alterations with Modified Single
TITLE OF INVENTION: Stranded Oligonucleotides
FILE REFERENCE: Napro-4
 NUMBER OF SEQ ID NOS: 4385
SOFTWARE: Friedman macro Napro4
 PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/192,179
 CURRENT APPLICATION NUMBER: US/09/818,875
CURRENT FILING DATE: 2001-03-27
 APPLICANT: Kmiec, Eric B.
APPLICANT: Gamper, Howar
APPLICANT: Rice, Michael
 ENERAL INFORMATION:
 FILE REFERENCE: MBHB00-831-F (400/009)
CURRENT APPLICATION NUMBER: US/09/825,805
CURRENT FILING DATE: 2001-09-27
 SOFTWARE: PatentIn version 3.0 EQ ID NO 782
 PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: 60/064,866
PRIOR FILING DATE: 1997-11-05
NUMBER OF SEQ ID NOS: 1558
 PRIOR APPLICATION NUMBER: 09/476,387
PRIOR FILING DATE: 1999-12-30
 PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER: 09/578,223
 TITLE OF INVENTION: Nucleotide Triphosphate and their Incorporation into Oligonucleo
FILE REFERENCE: MBHB00-831-F (400/009)
 ENGTH:
 , Match 0.9%; Local Similarity 62.5%; nes 10; Conservative
 2317 CATCAGAGTGATGTCT 2332
 1880 AGATGATGAAGATGAT 1895
 14;
 N
 Similarity
 CÁCCAGAGUGAUGUGU 17
AGATGATTCAGATGAT 16
 Conservative
 2000-05-23
 Howard B.
 0.9%;
 2001-09-27
 ; Score 12.8; D
; Pred. No. 1.7e
0; Mismatches
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 1.7e+02;
 Length 17;
 Length 17;
 0
 0
 Gaps
 Gaps
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 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-818-875-3314
 US-09-818-875-3314
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 ; ORGANISM: Homo sapiens US-09-818-875-1760
 RESULT 184
US-09-818-875-1760/c
 FILE REFERENCE: Napro-4
CURRENT APPLICATION NUMBER: US/09/818,875
CURRENT FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR PPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
PRIOR PILING DATE: 2000-03-27
PRIOR PILING DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR FILING DATE: 2000-10-30
PRIOR FILING DATE: 2000-10-30
 SOFTWARE: Friedman macro Napro4
SEQ ID NO 3314
LENGTH: 17
 Query Match 0.9%;
Best Local Similarity 87.5%;
Matches 14; Conservative
 SEQ ID NO 1760
 Sequence 1760, Application US/09818875 Publication No. US20030051270A1
 GENERAL INFORMATION:
 Sequence 3314, Application US/09818875 Publication No. US20030051270A1
 Matches
 Query Match
 GENERAL INFORMATION:
 APPLICANT: Kmiec, Eric B.
APPLICANT: Gamper, Howard B.
APPLICANT: Rice, Michael C.
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single
TITLE OF INVENTION: Stranded Oligonucleotides
 PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
 NUMBER OF SEQ ID NOS: 4385
 NUMBER OF SEQ ID NOS: 4385
SOFTWARE: Friedman macro Napro4
 CURRENT FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
 FILE REFERENCE: Napro-4
CURRENT APPLICATION NUMBER: US/09/818,875
CURRENT FILING DATE: 2001-03-27
 PRIOR FILING DATE: 2000-10-30
 TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single TITLE OF INVENTION: Stranded Oligonucleotides
 APPLICANT:
 APPLICANT: Kmiec, Eric B.
 PRIOR APPLICATION NUMBER: US 60/244,989
 APPLICANT:
 TYPE: DNA
 LENGTH: 17
 Local Similarity
1686 CCCAAAATGGGAGTTT 1701
 1880 AGATGATGAAGATGAT 1895
 17
 14;
 Gamper, Howard B. Rice, Michael C.
 Conservative
 0.9%;
 0
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 2;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 Length 17;
 Length 17;
 0,
 <u>.</u>
 Gaps
 0
 0
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 RESULT 187
US-09-780-533A-843/c
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 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-818-875-3315
 RESULT 186
US-09-818-875-3315/c
 밁
 ; TYPE: RNA; Homo sapiens US-09-780-533A-843
 PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR PRIOR PRIOR DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 4385
 APPLICANT: McSwiggen, Jim
APPLICANT: Chowrira, Bharat
APPLICANT: Chowrira, Bharat
APPLICANT: Habberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: PatentIn version 3.0
SEQ ID NO 843
 Publication No. US20 GENERAL INFORMATION:
 SOFTWARE: Friedman macro Napro4
SEQ ID NO 3315
LENGTH: 17
 Sequence 3315, Application US/09818875 Publication No. US20030051270A1
 Sequence 843, Application US/09780533A Publication No. US20030060611A1
 Matches
 Query Match
Best Local Similarity
 Matches
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 APPLICANT: Gamper, Howard B.
APPLICANT: Rice, Michael C.
TITLE OF INVENTION: Stranded Chromosomal Genomic Alterations with Modified Single
TITLE OF INVENTION: Stranded Oligonucleotides
FILE REFERENCE: Napro-4
 CURRENT APPLICATION NUMBER: US/09/818,875
CURRENT FILING DATE: 2001-03-27
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Blatt, Larry
 APPLICANT: Kmiec, Eric B. APPLICANT: Gamper, Howar
 ENGTH: 17
 1651 CTGGCAGGGGTCTCCG 1666
 1686 CCCAAAATGGGAGTTT 1701
17 CCGGCAGGGGTCCCCG 2
 17 CCCAAATTGGGTGTTT 2
 14;
 CCCAAATTGGGTGTTT 16
 0.9%;
ilarity 87.5%;
Conservative
 Conservative
 0.9%;
 ..
..
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8; DB 1;
Pred. No. 1:7e+02;
 Mismatches
 Mismatches
 DB 1;
 1;
 Length 17;
 Length 17;
 Indels
 0
 0
 Gaps
 Gaps
 0;
 0
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밁
 Query Match
Best Local Similarity
Matches 14; Conserva
 ; TYPE: RNA; Homo sapiens US-09-780-533A-2092
 RESULT 189
US-09-780-533A-2092/c
 밁
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 ; TYPE: RNA; Homo sapiens US-09-780-533A-845
 US-09-780-533A-845/c
 Sequence 845, Application US/09780533A Publication No. US20030060611A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
Matches 14; Conserv
 GENERAL INFORMATION:
 SEQ ID NO 2092
 Sequence 2092, Application US/09780533A Publication No. US20030060611A1
 SEQ ID NO 845
 APPLICANT: Ribozyne Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Chowrira, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHBOO, 878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR FILING DATE: 2001-02-11
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: Pstentin version 3.0
 APPLICANT: McSwiggen, Jim
APPLICANT: Chowrira, Bharat
APPLICANT: Chowrira, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00, 878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR PILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: Patentin version 3.0
 APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
APPLICANT: Chowrira, Bharat
 LENGTH:
 LENGTH: 17
 1572 GTCCAGCTCCTCCATG 1587
 1650 GCTGGCAGGGGTCTCC 1665
16 GrecaGGrerrecard 1
 16 GCCGGCAGGGGTCCCC 1
 Conservative
 Conservative
 0.9%;
 0.9%;
 0;
 0
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 2;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 <u>.</u>
 Length 17
 Length 17;
 0
 0
 Gaps
 0
 0
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RESULT 190 US-09-780-533A-2103/c

Sequence 2103, Application US/09780533A Publication No. US20030060611A1 GENERAL INFORMATION:

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US-09-877-478-2463
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 RESULT 192
 RESULT 191
US-09-780-533A-2341/c
 US-09-780-533A-2341
 US-09-780-533A-2103
 APPLICANT: McSwiggen, Jim
APPLICANT: Chowrira, Bhatat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene-
FILE REFERENCE: MBHB00, 878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: Patentin version 3.0
LENCTU. 17
 Publication No. US20 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 2341
 Sequence 2463, Application US/09877478
 Query Match
Best Local Similarity
 Sequence 2341, Application US/09780533A Publication No. US20030060611A1
 Query Match
Best Local
 APPLICANT:
APPLICANT:
 GENERAL INFORMATION:
 Matches
 APPLICANT: McSwiggen, Jim
APPLICANT: Chowrira, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NGGO Gene
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
 APPLICANT: Ribozyme Pharmaceuticals, APPLICANT: Draper, Kenneth
APPLICANT: McSwiggen, Ĵim
APPLICANT: Morrissey, Dave
TITLE OF INVENTION: Method and Reagent for Inhibiting Hepatitis B Virus Replication
FILE REFERENCE: MBHB00-845-H (400/029)
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Blatt, Larry
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 APPLICANT: Ribozyme Pharmaceuticals,
 Local Similarity
 1572 GTCCAGCTCCTCCATG 1587
 1653 GGCAGGGGTCTCCGAG 1668
 17
 17
 14;
 Draper, Kenne
Blatt, Larry
 Conservative
 Conservative
 US20030068301A1
 Larry
 0.9%;
 0.9%;
87.5%;
 N
 0
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 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8;
Pred. No. 1.
 Mismatches
 Mismatches
 1.7e+02
 DB 1;
 Length 17;
 Length 17;
 Indels
 0;
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 Gaps
 Gaps
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US-09-848-754A-326
 RESULT 194
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 ; TYPE: RNA; ORGANISM: Homo sapiens US-09-848-754A-184
 US-09-848-754A-184/c
 RESULT 193
 뭉
 US-09-877-478-2463
 Sequence 326, Application US/09848754A
Publication No. US20030073207A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MEHB00-958-I (400/018)
 CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
SOFTWARE: PatentIn version 3.0
SEQ ID NO 184
LENGTH: 17
 Sequence 184, Application US/09848754A
Publication No. US20030073207A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-I (400/018)
 Best Local Similarity Matches 14; Conserv
 Best Local Similarity Matches 11; Conserv
 Query Match
Best Local
 SOFTWARE: Pate
SEQ ID NO 2463
 Query Match
 PRIOR FILING DATE: 1994-02-07
PRIOR APPLICATION NUMBER: US 08/433,993
PRIOR FILING DATE: 1995-05-04
PRIOR APPLICATION NUMBER: US 08/434,504
PRIOR FILING DATE: 1995-05-04
PRIOR APPLICATION NUMBER: US 09/436,430
PRIOR APPLICATION NUMBER: US 09/436,430
 NUMBER OF SEQ ID NOS: 6586
 PRIOR FILING DATE: 1999-11-08
 PRIOR APPLICATION NUMBER: US 09/696,347
PRIOR FILING DATE: 2000-10-24
PRIOR APPLICATION NUMBER: US 08/193,627
 PRIOR APPLICATION NUMBER: US 09/531,025
PRIOR FILING DATE: 2000-03-20
PRIOR APPLICATION NUMBER: US 09/636,385
 PRIOR APPLICATION NUMBER: US 07/882,712 PRIOR FILING DATE: 1992-05-14
 CURRENT APPLICATION NUMBER: US/09/877,478
CURRENT FILING DATE: 2001-12-31
 PRIOR FILING DATE: 2000-08-09
 TYPE: RNA
ORGANISM: Hepatitis B virus
 ENGTH: 17
 1358 СССТССААСАСААА 1373
 2192 TGAAAATAGCAGACTT 2207
 16
 1 UGAAAACAGGAGACUU 16
 PatentIn version 3.0
 Conservative
 Conservative
NUMBER: US/09/848,754A
 0.9%;
 68.8%;
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 Score 12.8;
Pred. No. 1.
 Score 12.8;
Pred. No. 1
 Mismatches
 Mismatches
 1.7e+02;
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 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
US-09-848-754A-1167
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 US-09-848-754A-1481
 RESULT 196
 US-09-848-754A-1167/c
 RESULT 195
 US-09-848-754A-326
 Sequence 1481, Application US/09848754A

Publication No. US20030073207A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases of TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors

FILE REFERENCE: MBH800-958-T (400/018)

CURRENT APPLICATION NUMBER: US/09/848,754A

CURRENT FILING DATE: 2001-05-03

NUMBER OF SEQ ID NOS: 9645

SOFTWARE: Patentin version 3.0

SEQ ID NO 1481
 GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases

TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 1167
 NUMBER OF SEQ ID NOS: 9645
SOFTWARE: PatentIn version 3.0
SEQ ID NO 326
 Sequence 1167, Application US/09848754A
Publication No. US20030073207A1
 Matches
 Matches
 Query Match
 Query Match
 Query Match
 TYPE: RNA
ORGANISM: Homo sapiens
-09-848-754A-1481
 TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors FILE REFERENCE: MBHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
 CURRENT FILING DATE:
 TYPE: RNA
ORGANISM: Homo sapiens
 ENGTH:
 ENGTH: 17
 Local
 Local
 Local Similarity
 1358 CGCCTGGAAGAGAAA 1373
 2323 AGTGATGTCTGGTCCT 2338
 2314 ACTCATCAGAGTGATG 2329
 17
 14;
 1 AGUGAUGUCUGGAGCU 16
 17
N
 l Similarity
 Similarity 9; Conserv
 CGACTGCAAGAGAAAA 2
ACCCACCAGAGUGAUG 17
 Conservative
 Conservative
 Conservative
 NOS: 9645
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 0.9%;
 0.9%;
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 Score 12.8; Di
Pred. No. 1.7e
0; Mismatches
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 Score 12.8; D
Pred. No. 1.7e
2; Mismatches
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 1.7e+02;
 1.7e+02
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 DB 1;
 2
 Length 17;
 Length 17;
 Length 17;
 Indels
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 or Conditions
 or Conditions Relate
 Gaps
 Gaps
 Gaps
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 Relate
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 US-09-848-754A-3108/c
; Sequence 3108, Application US/09848754A
; Publication No. US20030073207A1
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 RESULT 197
US-09-848-754A-2894
 US-09-848-754A-3109/c
 RESULT 199
 US-09-848-754A-3108
 US-09-848-754A-2894
 CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
SOFTWARE: PATENTIN VEXISION 3.0
SEQ ID NO 2894
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate-
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
 Sequence 2894, Application US/09848754A Publication No. US20030073207A1 GENERAL INFORMATION:
 GENERAL INFORMATION:
 SEQ ID NO 3108
LENGTH: 17
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 3109 LENGTH: 17
 Sequence:3109, Application US/09848754A Publication No. US20030073207A1
 Matches
 Query Match
 Matches
 Query Match
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid
TITLE OF INVENTION: Levels of Epidermal G
FILE REFERENCE: MBHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-I (400/018)
 NUMBER OF SEQ ID NOS: 9645
SOFTWARE: PatentIn version
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 TYPE: RNA
ORGANISM: Homo sapiens
TYPE: RNA
 Local Similarity es 10; Conserv
 Local Similarity
 2664 TTTTTCTCCAGACCCC 2679
 2110 GGCATGGAGTACTTGG 2125
 17
 14;
 N
 GGCAUGAACUACUUGG 17
 TTTTCCTCCAGAGCCC
 Conservative
 Conservative
 version 3.0
 Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate-
Levels of Epidermal Growth Factor Receptors
 0.9%;
 0.9%;
 <u>.</u>.
 4
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8;
Pred. No. 1
 Mismatches
 Mismatches
 .7e+02
 DB 1;
 Length 17;
 Length 17;
 Indels
 Indels
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 0,
 Gaps
 0,
 . ;
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RESULT 201
US-09-776-474-679/c
 á
 문
 ; OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid
US-09-776-474-76
 ; ORGANISM: Homo sapiens
US-09-848-754A-3109
 NUMBER OF SEQ ID NOS: 2992
SOFTWARE: PatentIn version 3.0
SEQ ID NO 76
LENGTH: 17
 SEQ ID NO 679
 Matches 14; Conservative
 Matches 10;
 Query Match
 Query Match
 GENERAL INFORMATION:
 APPLICANT: Holman, Patricia
APPLICANT: Fattaey, Ali
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Inhibition of Checkpoint Kinase-1 (CHK
TITLE OF INVENTION: Enzyme
FILE REFERENCE: MBHB00-955-A (400/008)
CURRENT APPLICATION NUMBER: US/09/776,474
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR FILING DATE: 2000-03-02
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Inhibition of Checkpoint Kinase-1 (CHR
TITLE OF INVENTION: Enzyme
FILE REFERENCE: MBHB00-955-A (400/008)
CURRENT APPLICATION NUMBER: US/09/776,474
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR FILING DATE: 2000-03-02
 APPLICANT:
 SENERAL INFORMATION:
 ublication No.
 NUMBER OF SEQ ID NOS: 2992
SOFTWARE: PatentIn version 3.0
 APPLICANT: Ribozyme Pharmaceuticals, APPLICANT: Jarvis, Thale APPLICANT: Booher, Robert
 APPLICANT: Ribozyme Pharmaceuticals, APPLICANT: Jarvis; Thale
 APPLICANT:
 APPLICANT:
 LENGTH: 17
TYPE: RNA
 ORGANISM: Artificial Sequence
 TYPE: RNA
ORGANISM: Artificial Sequence
 FEATURE:
 Local Similarity
 Local Similarity
 2194 AAAATAGCAGACTTTG 2209
 2664 TTTTTCTCCAGACCCC 2679
 16
 N
 AAAAUCUCAGACUUUG 17
 Holman, Patricia
 Booher, Robert
 Fattaey, Ali
 Application US/09776474
o. US20030087847A1
 Application US/09776474
 Conservative
 Robert
 0.9%;
 0.9%;
 0;
 Score 12.8; D
Pred. No. 1.7e
0; Mismatches
 4; Mismatches
 Score 12.8;
Pred. No. 1.
 .7e+02
 .7e+02;
 DB
 B
 Length 17;
 Length 17;
 Indels
 0
 Gaps
 Gaps
 0
 0
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 ; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo Sapiens
US-09-930-423-386
 US-09-930-423-386/c
 US-09-776-474-1087
 RESULT 202
US-09-776-474-1087
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 ; FEATURE: ; OTHER INFORMATION: Description of Artificial Sequence: US-09-776-474-679
 NUMBER OF SEQ ID NOS: 4553
SOFTWARE: Patentin version 3.0
SEQ ID NO 386
LENGTH: 17
 GENERAL INFORMATION:
 Sequence 386, Application US/09930423 Publication No. US20030092003A1
 Matches
 Query Match
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 1087 LENGTH: 17
 Query Match
 GENERAL INFORMATION:
 Sequence 1087, Application US/09776474 Publication No. US20030087847A1
 Matches
 Query Match
 Matches
 APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERSNCE: MEHBOO, 918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930, 423
CURRENT FILING DATE: 2001-08-15
 APPLICANT: Fattaey, Ali
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Inhibition of Checkpoint Kinase-1 (CHK
TITLE OF INVENTION: Enzyme
FILE REFERENCE: MBHB00-955-A (400/008)
CURRENT APPLICATION NUMBER: US/99/776,474
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,983
PRIOR FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 2992
 APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
 APPLICANT: Ribozyme APPLICANT: Jarvis,
 APPLICANT:
 TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
 OTHER INFORMATION: Description of Artificial Sequence:
 0.9%;
Local Similarity 62.5%;
les 10; Conservation
 Local Similarity
nes 14; Conserv
 Match 0.9%; Local Similarity 87.5%;
 2195 AAATAGCAGACTTTGG 2210
 2163 AAATGTTTTGGTAACA 2178
 17
 AAAUCUCAGACUUUGG 16
 AAATGTTTTGATAAAA 2
 Holman, Patricia
 Conservative
 Conservative
 Pharmaceuticals, Inc
 Robert
 0.9%;
0; Mismatches
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8; DB 1
Pred. No. 1.7e+02
 Mismatches
 Mismatches
 DB 1;
 Length 17;
 Indels
 Indels
 Indels
 Nucleic
 Nucleic
0;
 <u>,,</u>
 0
 Acid
 Acid
Gaps
 Gaps
 Gaps
0,
 0,
 0
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1720 CTGGGCAAGCCCCTGG

1735

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US-09-780-164-225/c
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 RESULT 205
US-09-930-423-1169/c
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 S
 ; TYPE: RNA; Homo Sapiens US-09-930-423-660
 US-09-930-423-660
 밁
 RESULT 206
 ; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo Sapiens
US-09-930-423-1169
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: MBHB00,918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930,423
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARE: Patentin version 3.0
SEQ ID NO 1169
 Sequence 225, Application US/09780164 Publication No. US20030092646A1 GENERAL INFORMATION:
 GENERAL INFORMATION:
 Sequence 1169, Application US/09930423 Publication No. US20030092003A1
 SEQ ID NO 660
 GENERAL INFORMATION:
 Query Match
Best Local
 Matches 11; Conservative
 Sequence 660,
 Matches
 Query Match
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: MBHB00,918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930,423
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARE: PatentIn version 3.0
 Publication No.
 APPLICANT: Ribozyme Pharmaceuticals, APPLICANT: Blatt, Larry APPLICANT: McSwiggen, Jim
PPLICANT: McSwiggen, Jim
 LENGTH: 17
 Match 0.9%;
Local Similarity 87.5%;
 Match 0.9%;
Local Similarity 68.8%;
 1721 TGGGCAAGCCCCTGGG 1736
 2025 GGAGTACTCCTATGAC 2040
 17
 16
 14; Conservative
 N
 reseccascccccess
 CTGGGCCAGCCCCCGG 1
 Application US/09930423
o. US20030092003A1
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 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 Mismatches
 Inc
 Length 17;
 Length 17;
 Indels
 Indels
 0,
 0
 Gaps
 Gaps
 0
 0
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SOFTWARE: PatentIn version 3.0 SEQ ID NO 594

ORGANISM: Homo sapiens

TYPE: RNA LENGTH:

17

NUMBER OF SEQ ID NOS: 2603

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밁
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 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-780-164-225
 RESULT 208
US-09-780-164-594/c
 US-09-780-164-570
 RESULT 207
US-09-780-164-570
 CURRENT APPLICATION NUMBER: US/09/780,164
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/185,516
PRIOR FILING DATE: 2000-02-28
NUMBER OF SEQ ID NOS: 2603
SOFTWARE: PATENTIN VEXISION 3.0
SEQ ID NO 570
 Sequence 570, Application US/09780164
Publication No. US20030092646A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Inhibition of CD20
FILE REFERENCE: 400/010
 GENERAL INFORMATION:
 Sequence 594, Application US/09780164 Publication No. US20030092646A1
 Matches
 Query Match
Best Local Similarity
 Matches
 Query Match 0.9%;
Best Local Similarity 87.5%;
 SEQ ID NO 225
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Inhibition of
FILE REFERENCE: 400/010
CURRENT APPLICATION NUMBER: US/09/780,164
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/185,516
PRIOR APPLICATION NUMBER: 60/185,516
PRIOR FILING DATE: 2000-02-28
 FILE REFERENCE: 400/010
CURRENT APPLICATION NUMBER: US/09/780,164
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/185,516
PRIOR FILING DATE: 2000-02-28
NUMBER OF SEQ ID NOS: 2603
 SOFTWARE: PatentIn version 3.0
 TYPE: RNA
ORGANISM: Homo sapiens
 LENGTH: 17
 ENGTH:
 1703 CAAGAGATAAGCTGAC 1718
 1362 TGGAAGAGAAAAGGAG 1377
 13;
 16 TGTAAGAGAAAGGAG 1
 14;
 1 CAAGAGACAUGCUGAC 16
 Conservative
 Conservative
 81.2%;
 0.9%;
 0
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8;
Pred. No. 1.
 Mismatches
 Mismatches
 1.7e+02;
 DB 1;
 Length 17;
 Length 17;
 Indels
 CD20
 0,
 0
 Gaps
 Gaps
 0
 0
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; TYPE: RNA; Homo sapiens US-09-827-395A-360
 US-09-827-395A-360/c
 밁
 გ
 ; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-827-395A-198
 RESULT 209
US-09-827-395A-198/c
 US-09-780-164-594
 PRIOR APPLICATION NUMBER: 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 2617
SOFTWARE: PatentIn version 3.0
SEQ ID NO 360
LENGTH: 17
 PRIOR APPLICATION NUMBER: 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 2617
SOFTWARE: PatentIn version 3.0
SEQ ID NO 198
Query Match 0.9%;
Best Local Similarity 87.5%;
Matches 14; Conservative
 GENERAL INFORMATION:
 Publication No.
 Matches
 Query Match 0.9%;
Best Local Similarity 87.5%;
 Sequence 198, Application US/09827395A Publication No. US20030113891A1
 Matches
 GENERAL INFORMATION:
 APPLICANT: Bharat Chowrita
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO and NOGO Receptor
FILE REFERENCE: Method and Reagent for the Inhibition of NOGO and NOGO Receptor
CURRENT APPLICATION NUMBER: US/09/827,395A
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: 09/780,533
PRIOR APPLICATION NUMBER: 09/780,533
PRIOR FILING DATE: 2001-02-09
 APPLICANT: Ribozyme Pharmaceuticals, APPLICANT: Lawrence Blatt APPLICANT: James McSwiggen
 APPLICANT: Bharat Chowriza
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO and NOGO Receptor
FILE REFERENCE: MBHB00-878-C (400/017)
CURRENT APPLICATION NUMBER: US/09/827,395A
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: 09/780,533
PRIOR APPLICATION NUMBER: 09/780,533
PRIOR FILING DATE: 2001-02-09
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Lawrence Blatt
 APPLICANT:
 Local Similarity
 1572 GTCCAGCTCCTCCATG 1587
 1362 TGGAAGAGAAAAGGAG 1377
 14; Conservative
 17
 16 GTCCAGCTGCTCCAGG 1
 14;
 TGTAAGAGAGAAGGAG 2
 Lawrence Blatt
James McSwiggen
 Application US/09827395A
o. US20030113891A1
 Conservative
 0.9%;
 0
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 2;
 Score 12.8;
Pred. No. 1.
 0
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 Mismatches
 1.7e+02
 DB
 Length 17;
 Length 17;
 Length
 Indels
 Indels
0
 0
 0
Gaps
 Gaps
 0
 0
 <u>,</u>
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밁
 S
 ; TYPE: RNA; Homo sapiens US-09-827-395A-691
 US-09-827-395A-691/c
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 US-09-827-395A-611
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 RESULT 212
 US-09-827-395A-611/c
 Query Match
Best Local S
Matches 14
 Sequence 691, Application US/09827395A Publication No. US20030113891A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0
SEQ ID NO 611
LENGTH: 17
 Query Match
Best Local
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 691
 APPLICANT: James McSwiggen
APPLICANT: Bharat Chowrira
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO and NOGO Receptor C FILE REFERENCE: MBHB00-878-C (400/017)
CURRENT APPLICATION NUMBER: US/09/827,395A
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: 09/780,533
PRIOR APPLICATION NUMBER: 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 2617
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Lawrence Blatt
APPLICANT: James McSwiggen
 Sequence 611, Application US/09827395A Publication No. US20030113891A1
 PRIOR APPLICATION NUMBER: 09/780,533
PRIOR FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 2617
 APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Lawrence Blatt
APPLICANT: James McSwiggen
 CURRENT APPLICATION NUMBER: US/09/827,395A CURRENT FILING DATE: 2001-04-05
 APPLICANT: James McSwiggen
APPLICANT: Bharat Chowrira
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO and NOGO Receptor G
FILE REFERENCE: MBHB00-878-C (400/017)
 TYPE: RNA
ORGANISM: Homo sapiens
 LENGTH:
 Local Similarity
 Local Similarity
 1496 ACTTCAGCAGCCAGCC 1511
 1573 TCCAGCTCCTCCATGA 1588
 1496
16
 17
 14;
 ACTTCAGCAGCCAGCC 1511
 Conservative
 Conservative
 0.9%;
 0.9%;
 <u>,</u>
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8;
Pred. No. 1.
 Mismatches
 DB 1;
 В
 Length 17;
 0
 0
 Gaps
 Gaps
 0
 0
```

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RESULT 215
US-09-740-332-653/c
 뭉
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 ; OTHER INFORMATION: oligonucleotide substrate US-09-740-332-481
 RESULT 214
US-09-740-332-481/c
 밁
 á
 US-09-740-332-99
 US-09-740-332-99/c
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions
TITLE OF INVENTION: Hepatitis C Virus Infection
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: PatentIn version 3.0
SEQ ID NO 481
LENGTH: 17
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
CURRENT FAPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: Patentin version 3.0
 Sequence 653, Application US/09740332
Publication No. US20030125270A1
 SEQ ID NO 99
 Sequence 99, Application US/09740332 Publication No. US20030125270A1
 Sequence 481, Appropriate Publication No.
 Matches 14;
 LENGTH: 17
TYDE: RNA
ORGANISM: artificial sequence
FEATURE:
NAME/KEY: misc_feature
LOCATION:
 Matches
 Query Match
 Query Match
 NAME/KEY: misc_feature LOCATION:
 TYPE: RNA
ORGANISM: artificial sequence
FEATURE:
 OTHER INFORMATION: oligonucleotide substrate
 Local Similarity
 Local Similarity
 2047 CGTGTTCCTGAGGAGC 2062
 2452 AACTGCACCAACGAAC 2467
 16
 16
 14;
 AACTCCACCAACGATC 1
 Application US/09740332
 Conservative
 Conservative
 US20030125270A1
 0.9%;
87.5%;
 0.9%;
 <u>,</u>
 0
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 Mismatches
 Length 17;
 Length 17
 Indels
 0
 0;
 Gaps
 Gaps
 0
 0
 Relate
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RESULT 217
US-09-740-332-3135
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 , OTHER INFORMATION: oligonucleotide substrate US-09-740-332-1421
 밁
 RESULT 216
US-09-740-332-1421/c
 US-09-740-332-653
 NUMBER OF SEQ ID NOS: 9704
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3135
LENGTH: 17
 Sequence 3135, Application US/09740332

Publication No. US20030125270A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment

TITLE OF INVENTION: Hepatitis C Virus Infection
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 1421 LENGTH: 17
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 653 LENGTH: 17
 Matches
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 Sequence 1421, Application US/09740332
Publication No. US20030125270A1
 Matches
 Query Match
Best Local Similarity
 TITLE OF INVENTION: Hepatitis C Virus Infection FILE REFERENCE: RPI 400/003 CURRENT APPLICATION NUMBER: US/09/740,332 CURRENT FILING DATE: 2001-03-26
 APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate.
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
FULL REFERENCE: RPI 400/003
FULL REFERENCE: US/09/740,332
FULL REFERENT APPLICATION NUMBER: US/09/740,332
FULL REFERENT APPLIAGE DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
 FILE REFERENCE: RPI 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
 ORGANISM: artificial sequence FEATURE:
 LOCATION:
OTHER INFORMATION: oligonucleotide substrate
 ORGANISM: artificial sequence FEATURE:
 TYPE: RNA
 NAME/KEY: misc_feature
 TYPE: RNA
 NAME/KEY: misc_feature
 LOCATION:
 1737
 1473 GAAGAACACGACCAAG 1488
RNA
 17 GAAGAACACGAGGAAG 2
 16
 14;
 14;
 AGAAGGTTGCTTTGGG 1752
 Conservative
 Conservative
 0.9%;
 0.9%;
 0
 °;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 Mismatches
 ٠,
 Length 17;
 Length 17
 Indels
 Diseases
 0
 0,
 or Conditions Relate
 Gaps
 Gaps
 0
 0
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ORGANISM: artificial sequence

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RESULT 219
US-09-740-332-4457
US-09-740-332-4457
Sequence 4457, Application US/09740332
Publication No. US20030125270A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Heparitis C Virus Infection
FILE REFERENCE: RPI 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
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 ; OTHER INFORMATION: oligonucleotide substrate US-09-740-332-3902
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 ; OTHER INFORMATION: oligonucleotide substrate US-09-740-332-4457
 US-09-740-332-3902
 RESULT 218
 US-09-740-332-3135
 SEQ ID NO 3902
LENGTH: 17
TYPE: RNA
 NUMBER OF SEQ ID NOS: 9704
SOFTWARE: PatentIn version 3.0
SEQ ID NO 4457
 Query Match 0.9%;
Best Local Similarity 75.0%;
Matches 12; Conservative
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
 Query Match 0.5
Best Local Similarity 62.5
Matches 10; Conservative
 Matches
 Sequence 3902, Application US/09740332 Publication No. US20030125270A1
 NUMBER OF SEQ ID NOS: 9704
SOFTWARE: PatentIn version 3.0
 CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
 FEATURE: misc_feature
 ORGANISM: artificial sequence
 TYPE: RNA
 NAME/KEY: misc_feature
 ORGANISM: artificial sequence FEATURE:
 OTHER INFORMATION: oligonucleotide substrate
 LOCATION:
 FEATURE: misc_feature
 OCATION:
 ENGTH: 17
 1737 AGAAGGTTGCTTTGGG 1752
 1473 GAAGAACACGACCAAG 1488
 14; Conservative
 N
 GAAGAACACGAGGAAG 17
 AGGAGGGUGCUUUGGG 16
 0.9%;
 62.5%;
 0;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
2; Mismatches 2;
 Score 12.8;
Pred. No. 1.
 4.
 Score 12.8;
Pred. No. 1
 Mismatches
 Mismatches
 8; DB 1;
1.7e+02;
 DB 1;
..7e+02;
 Length 17;
 Length 17;
 Length 17;
 Indels
 Indels
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 0
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Gaps
 Gaps
 Gaps
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 0
 0
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RESULT 222
US-09-745-237A-1169/c
US-09-745-237A-1169/c
; Sequence 1169, Application US/09745237A
; Publication No. US20030143708A1
; GENERAL INFORMATION:
GENERAL INFORMATION:
 밁
 ; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-745-237A-660
 RESULT 221
US-09-745-237A-660
 문
 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-745-237A-386
 US-09-745-237A-386/c
US-09-745-237A-386/c
; Sequence 386, Application US/09745237A
; Publication No. US20030143708A1
 밁
 5
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: MCSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: 400/007 (MBHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: Patentin version 3.0
SEQ ID NO 660
 0.9%;
pest Local Similarity 87.5%;
Matches 14; Conservation
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 386 LENGTH: 17
 Matches
 Query Match
 Sequence 660, Appropriate Publication No.
 GENERAL INFORMATION:
 GENERAL INFORMATION:
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: 400/007 (MBHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
 Local Similarity es 11; Conserv
 2025 GGAGTACTCCTATGAC 2040
 1720 CTGGGCAAGCCCCTGG 1735
 2454 CTGCACCAACGAACTG 2469
 16
 2 GGAGUACAACUAUGAC 17
 CTGGGCCAGCCCCCGG
 CUCCACCAACGAUCUG 16
 Application US/09745237A
o. US20030143708A1
 Conservative
 0.98;
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 Score 12.8;
Pred. No. 1.
 Score 12.8;
Pred. No. 1.
 Mismatches
 Mismatches
 1.7e+02;
ches 2;
 .7e+02;
 BB
 Length 17;
 Length
 Indels
 Indels
 0;
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 Gaps
 Gaps
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 0
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 US-09-792-818-283/c
 US-09-745-237A-1169
 US-10-238-700-661
SOFTWARE: Pat
SEQ ID NO 661
LENGTH; 17
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 1169
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 283
 Query Match
Best Local (
 GENERAL INFORMATION:
 Sequence 661, Application US/10238700 Publication No. US20030153521A1
 Query Match
Best Local
 Publication No. US20030134806A1 GENERAL INFORMATION:
 Matches
 ORGANISM: Homo sapiens
 PRIOR APPLICATION NUMBER: PCT/US 02/16840 PRIOR FILING DATE: 2002-05-29 PRIOR APPLICATION NUMBER: US 60/318,471 PRIOR FILING DATE: 2001-09-10 NUMBER OF SEQ ID NOS: 4666
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
 APPLICANT: Hamblin, Paul
APPLICANT: Ellis, Jonathan
TITLE OF INVENTION: Method and Reagent for the Inhibition of Grb-2-related with Inservation of Invention: (GRID) Gene
TITLE OF INVENTION: (GRID) Gene
FILE REFERENCE: MBHB00-901-A (400/013)
CURRENT FILING INDICATION NUMBER: US/09/792,818
CURRENT FILING DATE: 2001-02-23
NUMBER OF SEQ ID NOS: 2304
 FILE REFERENCE: 400/007 (MBHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
 APPLICANT:
 APPLICANT: Ribozyme Pharmaceuticals, APPLICANT: Jarvis, Thale
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 TYPE: RNA
 ENGTH: 17
 1869 GTCAGAGATGGAGATG 1884
 1721 TGGGCAAGCCCCTGGG 1736
 17 GACAGAGATGGAGAAG 2
 14;
 17 TGGGCCAGCCCCGGG 2
 Similarity
 Similarity
 PatentIn version 3.0
 Von Carlowitz, Ira
McSwiggen, Jim
Hamblin, Paul
 Application US/09792818
 Conservative
 Conservative
 0.9%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 1.7e+02;
 0.9%;
 Score 12.8; D
Pred. No. 1.7e
0; Mismatches
 <u>,</u>
 Mismatches
 DB 1;
 Length 17;
 Indels
 Indels
 0
 0
 0;
 0
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Sequence 3208, Application US/10238700
Publication No. US20030153521A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
ITILE OF INVENTION: Nucleic Acid Treatment of
FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT FILING DATE: 2002-09-18
PRIOR APPLICATION NUMBER: PCT/US 02/16840
PRIOR FILING DATE: 2002-09-18
PRIOR FILING DATE: 2002-09-19
PRIOR FILING DATE: 2001-09-10
INUMBER OF SEQ ID NOS: 4666
SOFTWARE: Patentin version 3.0
SEQ ID NO 3208
LENGTH: 17
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 US-10-238-700-3208
 RESULT 226
US-10-238-700-3208
 US-10-238-700-944
 APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of
FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
PRIOR APPLICATION NUMBER: PCT/US 02/16840
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2001-09-10
PRIOR FILING DATE: 2001-09-10
NUMBER OF SEQ ID NOS: 4666
 US-10-238-700-944
 RESULT 225
 US-10-238-700-661
 SEQ ID NO 944
 Sequence 944, Applic
Publication No. US20
GENERAL INFORMATION:
Best Local Similarity Matches 11; Conserv
 Matches
 Best Local Similarity
 Query Match
 Best Local Similarity
Matches 10; Conserv
 Query Match
 Query Match
 LENGTH: 17
TYPE: RNA
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, James
 SOFTWARE: PatentIn version 3.0
 TYPE: RNA
ORGANISM: Homo sapiens
 ORGANISM: Homo sapiens
 ORGANISM: Homo sapiens
 2151 TTTAGCAGCCAGAAAT 2166
 2190 GATGAAAATAGCAGAC 2205
 13;
 N
 2 UUUAGUAACCAGAAAU 17
 GAAGAAUAUAGCAGAC 17
 Application US/10238700
 Conservative
 Conservative
 Conservative
 US20030153521A1
 0.9%;
 68.8%;
 0.9%;
 4:
 Score 12.8;
Pred. No. 1.
 Score 12.8;
Pred. No. 1.
 Score 12.8;
Pred. No. 1.
 Mismatches
 Mismatches
 8; DB 1;
1.7e+02;
ches 2;
 1.7e+02;
 .7e+02
 of Diseases or Conditions Related to Level
 of Diseases or Conditions Related to Level
 DB 1;
 DB 1;
 Length 17;
 Length 17;
 Length 17;
 Indels
 0
 0
 <u>,,</u>
 Gaps
 Gaps
 Gaps
 0
 0
 0
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1512 GGCTGTGCACAAGCTG 1527

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RESULT 229
US-10-061-201-121/c
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 US-10-253-904-31
 US-10-253-904-31/c
 RESULT 228
 US-10-238-700-3544/c
 RESULT 227
 US-10-238-700-3544
Sequence 121, Application US/10061201 Publication No. US20030166229A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 31
LENGTH: 17
TYPE: DNA
ORGANISM: Artificial Sequence
 Matches
 Sequence 31, App
Publication No.
 Query Match
 Query Match
 SEQ ID NO 3544
 Sequence 3544, Application US/10238700 Publication No. US20030153521A1
 Matches
 GENERAL INFORMATION:
 GENERAL INFORMATION:
 CURRENT APPLICATION NUMBER: US/10/253,904
CURRENT FILING DATE: 2002-09-25
NUMBER OF SEQ ID NOS: 51
 APPLICANT: ALLIGNET, JEANINE
TITLE OF INVENTION: POLYNUCLEOTIDES AND THEIR USE FOR DETECTING RESISTANCE
TITLE OF INVENTION: TO STREPTOGRAMIN A OR TO STREPTOGRAMIN B AND RELATED
TITLE OF INVENTION: COMPOUNDS
FILE REFERENCE: 03715-0059
 FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
PRIOR APPLICATION NUMBER: PCT/US 02/16840
 PRIOR APPLICATION NUMBER: US 60/318,471 PRIOR FILING DATE: 2001-09-10 NUMBER OF SEQ ID NOS: 4666
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwijgen, James TITLE OF INVENTION: Nucleic Acid Treatment FILE REFERENCE: 400/057 (MBHB01-1118-A)
 APPLICANT: EL SOLH, NEVINE APPLICANT: ALLIGNET, JEAN:
 SOFTWARE: PatentIn version 3.0
 PRIOR FILING DATE: 2002-05-29
 LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
 OTHER INFORMATION: Description of Artificial Sequence: Primer
 FEATURE:
 Local
 Local
 1741 GGTTGCTTTGGGCAAG 1756
 2393 TTCCCGTGGAGGAACT 2408
 17
 17
 14;
 14; Conservative
 Similarity
 Similarity
 GGTTGCTTTTGCCAAG 2
 GGCUGUGCACAGACUG 16
 Application US/10253904
No. US20030158135A1
 Conservative
 0.9%;
 0.9%;
 <u>,</u>
 0,
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 2;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 of Diseases or Conditions Related to Leve
 Length 17;
 Length 17;
 Indels
 0
 0
 Gaps
 Gaps
 0
 0
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GENERAL INFORMATION:

US20030166229A1

TITLE OF INVENTION:

HUMAN POSH-LIKE PROTEIN 1

PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00669

FILE REFERENCE: PB0178
CURRENT APPLICATION NUMBER: US/10/061,201
CURRENT FILING DATE: 2002-01-30

PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 09/864,761
PRIOR APPLICATION NUMBER: US 60/328,205
PRIOR FILING DATE: 2001-10-10

Aeomica Sequence Listing Engine

ID NOS: 4162

PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00668

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US-10-061-201-124/c
 RESULT 230
 ; ORGANISM: Homo sapiens US-10-061-201-121
Sequence 124, A Publication No.
 Matches
 Query Match
Best Local Similarity
 SOFTWARE: Ae
SEQ ID NO 121
 PRIOR
PRIOR
 FILE REFERENCE: PB0178
CURRENT APPLICATION NUMBER: US/10/061,201
CURRENT FILING DATE: 2002-01-30
CURRENT FILING DATE: DOTT/HR01/00666
 PRIOR APPLICATION NUMBER: US 60/328,205
PRIOR FILING DATE: 2001-10-10
 NUMBER OF SEQ ID NOS: 4162
 PRIOR APPLICATION NUMBER: PCT/US01/00669
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
 TITLE OF INVENTION: HUMAN POSH-LIKE PROTEIN
 TYPE: DNA
 RIOR
 PRIOR APPLICATION NUMBER: PCT/US01/00665
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00664
 PRIOR APPLICATION NUMBER: PCT/US01/00667
 ENGTH:
 APPLICATION NUMBER: US 09/864,761
FILING DATE: 2001-05-23
 APPLICATION NUMBER: PCT/
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 1565 CGGCTGAGTCCAGCTC 1580
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663
 FILING DATE: 2001-01-30
 17 CTGCTGAGTTCAGCTC 2
 14;
 Aeomica Sequence Listing Engine
 Application US/10061201
 Conservative
 2001-01-30
 0.9%;
 PCT/US01/00670
 Score 12.8;
Pred. No. 1.
 Mismatches
 DB 1;
.7e+02;
 Length 17;
 0
 Gaps
```

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RESULT 232
US-09-817-879-481/c
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 RESULT 231
US-09-817-879-99/c
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 S
 US-09-817-879-481
 US-09-817-879-99
 US-10-061-201-124
 Sequence 481, Application US/09817879

Publication No. US20030171311A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate

TITLE OF INVENTION: Hepatitis C Virus Infection

FILE REFERENCE: MBH800-801-F

CURRENT APPLICATION NUMBER: US/09/817,879

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9703

SOFTWARE: PatentIn version 3.0

SEQ ID NO 481
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection

FILE REFERENCE: MEHBOO-801-F
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: Patentin version 3.0
SEQ ID NO 99
LENGTH: 17
 SEQ ID NO 124
LENGTH: 17
 Sequence 99, App
Publication No.
 Matches
 Matches
 Query Match
 Query Match
 Best Local Similarity
 Query Match
 LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
 NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: oligonucleotide substrate
 OTHER INFORMATION: oligonucleotide substrate
 FEATURE:
 ORGANISM: artificial sequence
 FEATURE:
 TYPE: RNA ORGANISM: artificial sequence
 NAME/KEY: misc_feature
 ENGTH: 17
y Match 0.9%;
Local Similarity 87.5%;
hes 14; Conservative
 Match 0.9%;
Local Similarity 87.5%;
 2452 AACTGCACCAACGAAC 2467
 1563 TTCGGCTGAGTCCAGC 1578
 14; Conservative
 16
 16 AACTCCACCAACGATC 1
 14; Conservative
 Application US/09817879
 0.9%; Score 12.8; DB 1; 87.5%; Pred. No. 1.7e+02;
 0
 ٥,
 0; Mismatches
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 Mismatches
 Length 17;
 Length 17;
 Length 17;
 Indels
 Indels
 Indels
0
 0
 ٥,
Gaps
 Gaps
 Gaps
0
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 RESULT 233
US-09-817-879-653/c
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 S
 ; OTHER INFORMATION: oligonucleotide substrate US-09-817-879-1421
 RESULT 234
US-09-817-879-1421/c
 ; OTHER INFORMATION: oligonucleotide substrate US-09-817-879-653
 APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate.

TITLE OF INVENTION: Hepatitis C Virus Infection

FILE REFERENCE: MBHB00-801-F

CURRENT APPLICATION NUMBER: US/09/817,879

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9703

SOFTWARE: PatentIn version 3.0

SEQ ID NO 1421
 Sequence 1421, Application US/09817879 Publication No. US20030171311A1 GENERAL INFORMATION:
 Sequence 653, Applic
Publication No. US20
GENERAL INFORMATION:
 SEQ ID NO 653
LENGTH: 17
 Matches
 Query Match
Best Local Similarity
 Matches
 Query Match
Best Local Similarity
 CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: Patentin version 3.0
 APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate.
TITLE OF INVENTION: Hepatitis C Virus Infection
 TITLE OF INVENTION: Hepatitis C Virus Infection FILE REFERENCE: MBHB00-801-F
 FEATURE:
NAME/KEY: misc_feature
LOCATION:
 LENGTH: 17
 FEATURE: misc_feature
 ORGANISM: artificial sequence
 ORGANISM: artificial sequence
 TYPE: RNA
 LOCATION:
 1737
 2047
 1473 GAAGAACACGACCAAG 1488
 14;
 17 GAAGAACACGAGGAAG 2
 16
16
 14;
 AGAAGGTTGCTTTGGG 1752
 CGTGTTCCTGAGGAGC 2062
 Application US/09817879
o. US20030171311A1
 Conservative
 Conservative
 0.9%;
 0.9%;
 0;
 0;
 Score 12.8;
Pred. No. 1
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
 Mismatches
 1.7e+02;
 DB 1;
 DB 1;
 Length 17;
 Length 17
 Indels
 0
 ٥,
 Gaps
 Gaps
 0
 0
```

RESULT 235 US-09-817-879-3135

Sequence 3135,

Application US/09817879

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RESULT 237
US-09-817-879-4457
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 5
 US-09-817-879-3902
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 ફ
 US-09-817-879-3902
 RESULT 236
 ; OTHER INFORMATION: oligonucleotide substrate US-09-817-879-3135
 Sequence 3902, Application US/09817879

Publication No. US20030171311A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate

TITLE OF INVENTION: Hepatitis C Virus Infection

FILE REFERENCE: MBHB00-801-F

CURRENT APPLICATION NUMBER: US/09/817,879

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9703

SOFTWARE: PatentIn version 3.0
Sequence 4457, Application US/09817879
Publication No. US20030171311A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MBHB00-801-F
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
 Query Match
Best Local Similarity
Matches 14; Conserv
 SOFTWARE: PatentIn version 3.0
SEQ ID NO 3135
LENGTH: 17
TYPE: RNA
 SEQ ID NO 3902
 GENERAL INFORMATION:
 Query Match
Best Local
 Matches
 Publication No. US20030171311A1
 APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MBHB00-801-F
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
 CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
 FEATURE:
NAME/KEY: misc_feature
 OTHER INFORMATION: oligonucleotide substrate
 NAME/KEY: misc_feature
LOCATION:
 ORGANISM: artificial sequence
 TYPE: RNA
 FEATURE:
 ORGANISM: artificial sequence
 LOCATION:
 ENGTH: 17
 0.9%;
Local Similarity 62.5%;
les 10; Conservative
 1737 AGAAGGTTGCTTTGGG 1752
 1473 GAAGAACACGACCAAG 1488
 N
 GAAGAACACGAGGAAG 17
 AGGAGGGUGCUUUGGG 16
 Conservative
 0.9%;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8;
Pred. No. 1.
 Mismatches
 DB 1;
.7e+02;
 Length 17,
 Length 17;
 0
 0
 Gaps
 0
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US-10-339-793-443/c
 RESULT 239
 ; OTHER INFORMATION: Description of Artificial Sequence: primer US-10-392-970-29
 US-10-392-970-29/c
 US-09-817-879-4457
 Sequence 443, Application US/10339793
Publication No. US20030180764A1
GENERAL INFORMATION:
APPLICANT: Lynx Therapeutics, Inc.
APPLICANT: Shang, Jin
APPLICANT: Bowen, Benjamin
 CURRENT APPLICATION NUMBER: US/10/339,793
CURRENT FILING DATE: 2003-01-08
NUMBER OF SEQ ID NOS: 443
SOFTWARE: PatentIn version 3.1
SEQ ID NO 443
 Query Match
Best Local Similarity
Matches 14; Conser
 SEQ ID NO 29
 GENERAL INFORMATION:
 Sequence 29, Application US/10392970 Publication No. US20030176679A1
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 4457
 Matches
 Query Match
 TITLE OF INVENTION: GENES AFFECTED BY CHOLESTEROL TREATMENT AND DURING ADIPOGENESIS FILE REFERENCE: 37-000310US
 APPLICANT: Allignet, Jeanine
TITLE OF INVENTION: POLYNUCLEOTIDES AND THEIR USE FOR DETECTING RESISTANCE
TITLE OF INVENTION: TO STREPTOGRAMIN A OR TO STREPTOGRAMIN B AND RELATED
TITLE OF INVENTION: COMPOUNDS
TILE REFERENCE: 03495.0173-00000
CURRENT APPLICATION NUMBER: US/10/392,970
CURRENT FILLING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: US/09/099,932
PRIOR FILLING DATE: 1998-06-19
PRIOR FILLING DATE: 1998-06-19
 NUMBER OF SEQ ID NOS: 50 SOFTWARE: Patentin Ver. 2.0
 PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/050,380 PRIOR FILING DATE: EARLIER FILING DATE: 1997-06-20
 APPLICANT: El Solh, APPLICANT: Alligne
 LENGTH: 17
TYPE: DNA
ORCANISM: Artificial Sequence
FEATURE:
 ORGANISM: artificial sequence FEATURE: misc_feature
 TYPE: RNA
TYPE: DNA
 OTHER INFORMATION: oligonucleotide substrate
 LENGTH:
 LOCATION:
 Local Similarity
 1741 GGTTGCTTTGGGCAAG 1756
 2454 CTGCACCAACGAACTG
 17
 μ
 GGTTGCTTTTGCCAAG
 CUCCACCAACGAUCUG 16
 Conservative
 Conservative
 Nevine
 0.9%;
 0.9%;
 N
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 2;
 Score 12.8; D
Pred. No. 1.7e
0; Mismatches
 Score 12.8;
Pred. No. 1.
 Mismatches
 1.7e+02
 ; DB 1;
1.7e+02;
 B
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 Length 17
 Length 17;
 Indels
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 Gaps
 Gaps
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 RESULT 240
US-10-209-787-1759
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 US-10-339-793-443
 US-10-209-787-1760/c
 RESULT 241
 PRIOR APPLICATION NUMBER: US 09/818,875
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-06-01
 Query Match 0.9%;
Best Local Similarity 87.5%;
Matches 14; Conservative
 Sequence 1760, Application US/10209787 Publication No. US20030217377A1 GENERAL INFORMATION:
 SEQ ID NO 1759
LENGTH: 17
 Query Match
Best Local
 GENERAL INFORMATION
 Sequence 1759, Application US/10209787 Publication No. US20030217377A1
 Matches
 Publication No.
 ORGANISM: Homo sapiens
-10-209-787-1759
 APPLICANT: Kmiec, Eric B.

APPLICANT: Gamper, Howard B.

APPLICANT: Rice, Michael C.

TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single

TITLE OF INVENTION: Stranded Oligonucleotides
 PRIOR APPLICATION NUMBER: US 09/818,875
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR FILING DATE: 2000-03-27
PRIOR FILING DATE: 2000-03-27
 FILE REFERENCE: Napro-4
CURRENT APPLICATION NUMBER: US/10/209,787
CURRENT FILING DATE: 2002-07-30
 PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR FILING DATE: 2000-10-30
 CURRENT APPLICATION NUMBER: US/10/209,787
CURRENT FILING DATE: 2002-07-30
 TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single TITLE OF INVENTION: Stranded Oligonucleotides
 APPLICANT: Kmiec, Eric B.
APPLICANT: Gamper, Howar
APPLICANT: Rice, Michael
 FILE REFERENCE: Napro-4
 SOFTWARE: Friedman macro Napro4
 ORGANISM: Homo sapiens
 TYPE: DNA
 Local
 1880 AGATGATGAAGATGAT 1895
 2238 CTATTACAAAAAGACC 2253
 FILING DATE: 2000-10-30 OF SEQ ID NOS: 4385
 16
 ch 0.9%; l Similarity 87.5%; 14; Conservative
 CTTTTACAAAAAGATC 1
CATION NUMBER: US 60/208,538
 AGATGATTCAGATGAT 16
 Gamper, Howard B. Rice, Michael C.
 Pred. No. 1.7e
0; Mismatches
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 2;
 1.7e+02;
 DB
 1;
 Length 17;
 Length 17;
 Indels
 0
 ٥,
 Gaps
 0
 0
```

US-10-209-787-3315/c

Sequence 3315, Application US/10209787 Publication No. US20030217377A1 GENERAL INFORMATION:

APPLICANT: Kmiec, Eric B.
APPLICANT: Gamper, Howar
APPLICANT: Rice, Michael

Gamper, Howard B. Rice, Michael C.

FILE REFERENCE: Napro-4
CURRENT APPLICATION NUMBER: US/10/209,787
CURRENT FILING DATE: 2002-07-30

PRIOR APPLICATION NUMBER: US 09/818,875

TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single TITLE OF INVENTION: Stranded Oligonucleotides

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RESULT 243
 ; ORGANISM: Homo sapiens US-10-209-787-3314
 RESULT 242
US-10-209-787-3314
 US-10-209-787-1760
 PRIOR APPLICATION NUMBER: US 09/818,875
PRIOR FILLING DATE: 2001-03-27
PRIOR PELLING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR APPLICATION NUMBER: US 60/192,179
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR PRILING DATE: 2000-10-30
PRIOR FILLING DATE: 2000-10-30
PRIOR FILLING DATE: 2000-10-30
PRIOR FILLING DATE: 2000-10-30
 SOFTWARE: Friedman macro Napro4
SEQ ID NO 1760
 SOFTWARE: Friedman macro Napro4
SEQ ID NO 3314
 Publication No. US20 GENERAL INFORMATION:
 Sequence 3314, Application US/10209787 Publication No. US20030217377A1
 Matches
 Query Match
Best Local Similarity
 Best Local Similarity Matches 14; Conserv
 Query Match
 PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 4385
 APPLICANT: Gamper, Howard B.
APPLICANT: Rice, Michael C.
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single
TITLE OF INVENTION: Stranded Oligonucleotides
 FILE REFERENCE: Napro-4
CURRENT APPLICATION NUMBER: US/10/209,787
CURRENT FILING DATE: 2002-07-30
 APPLICANT: Kmiec, Eric B.
APPLICANT: Gamper, Howar
 TYPE: DNA
ORGANISM: Homo sapiens
 TYPE: DNA
 LENGTH: 17
 ENGTH:
 1686 CCCAAAATGGGAGTTT 1701
 1880 AGATGATGAAGATGAT 1895
 17
 14;
 1 CCCAAATTGGGTGTTT 16
 SEQ ID NOS: 4385
 Conservative
 Conservative
 0.9%;
87.5%;
 0.9%;
 <u>,</u>
 0,
 Score 12.8;
Pred. No. 1
 Score 12.8;
Pred. No. 1
 Mismatches
 Mismatches
 1.7e+02;
 L.7e+02;
 BB
 DB 1;
 Length 17;
 Length 17;
 Indels
 Indels
 0
 <u>,</u>
 Gaps
 0
 0
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RESULT 245
US-10-060-830-701/c
; Sequence 701, Application US/10060830
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 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-060-830-700
 US-10-060-830-700/c
 RESULT 244
 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-209-787-3315
 Matches
 SEQ ID NO 3315
 Matches 14; Conservative
 Query Match
 Sequence 700, A
Publication No.
 PRIOR APPLICATION NUMBER: US 60/208,538
PRIOR FILING DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 4385
SOFTWARE: Friedman macro Napro4
 SOFTWARE: Aeomica Sequence Listing Engine
 CURRENT APPLICATION NUMBER: US/10/060,830
CURRENT FILING DATE: 2002-01-30
 PRIOR FILING DATE: 2001-09-25
 PRIOR APPLICATION NUMBER: PCT/US01/00667
 FILE REFERENCE: PB0169
 APPLICANT: Gu, Yizhong
APPLICANT: Nguyen, Cu
 ENERAL INFORMATION
 PPLICANT: Nguyen, Cung-Tuong
ITLE OF INVENTION: HUMAN LCCL DOMAN CONTAINING PROTEIN
 Local Similarity
 Match 0.9%;
Local Similarity 87.5%;
 APPLICATION NUMBER: US 09/864,761
FILING DATE: 2001-05-23
APPLICATION NUMBER: US 60/325,062
 APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: PCT/US01/00664 FILING DATE: 2001-01-30
 FILING DATE:
 APPLICATION NUMBER: US 60/192,179 FILING DATE: 2000-03-27
 FILING DATE: 2001-03
APPLICATION NUMBER:
FILING DATE: 2000-03
 2054 CTGAGGAGCAGATGAC 2069
 1686 CCCAAAATGGGAGTTT 1701
 LING DATE: 2001-01-30
 CTGAGGAGCAGCTGTC 2
 Application US/10060830 to. US20030032154A1
 Conservative
 2001-01-30
 2001-03-27
 2000-03-
 0.9%;
 2002-01-30
 US 60/192,176
3-27
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 Score 12.8;
Pred. No. 1
 Score 12.8;
Pred. No. 1.
 1.7e+02
 DB
 Length 17;
 Length 17;
 Indels
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 0
 Gaps
 Gaps
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 US-10-060-756A-144
 US-10-060-830-701
 SOFTWARE: Ae
SEQ ID NO 144
 Sequence 144, Application US/10060756A Publication No. US20030046717A1 GENERAL INFORMATION:
 Matches
 Query Match
Best Local Similarity
 SEQ ID NO 701
 Publication No. US20030032154A1 GENERAL INFORMATION:
 NUMBER OF SEQ ID NOS: 4804
 PRIOR APPLICATION NUMBER: US 60/327,898 PRIOR FILING DATE: 2001-10-09
 PRIOR APPLICATION NUMBER: US 09/864,761 PRIOR FILING DATE: 2001-05-23
 CURRENT APPLICATION NUMBER: US/10/060,756A
CURRENT FILING DATE: 2002-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
 APPLICANT: Zhang, Ji
TITLE OF INVENTION:
 CURRENT APPLICATION NUMBER: US/10/060,830 CURRENT FILING DATE: 2002-01-30
 FILE REFERENCE: PB0169
 APPLICANT: Nguyen, Cung-Tuong
TITLE OF INVENTION: HUMAN LCCL DOMAN CONTAINING PROTEIN
 APPLICANT: Gu, Yizhong APPLICANT: Nguyen, Cu
 PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
 NUMBER OF SEQ ID NOS: 1123
 PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
 PRIOR APPLICATION NUMBER: PCT/US01/00664
 FILE REFERENCE: PB0177
 PRIOR FILING DATE:
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 09/864,761
 PRIOR APPLICATION NUMBER: PCT/US01/00669
 PRIOR APPLICATION NUMBER: PCT/US01/00664 PRIOR FILING DATE: 2001-01-30
 ORGANISM: Homo sapiens
 TYPE: DNA
 PRIOR APPLICATION NUMBER:
 RIOR FILING DATE:
 RIOR APPLICATION NUMBER: PCT/US01/00668
 RIOR APPLICATION NUMBER: PCT/US01/00665
ORGANISM: Homo sapiens
 LENGTH: 17
 RIOR FILING DATE: 2001-01-30
 APPLICATION NUMBER: US 60/325,062 FILING DATE: 2001-09-25
 2054 CTGAGGAGCAGATGAC 2069
 FILING DATE: 2001-05-23
 16 CTGAGGAGCAGCTGTC 1
 14;
 Aeomica Sequence Listing Engine
 Aeomica Sequence Listing Engine
 Conservative
 2001-01-30
 2001-01-30
 HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
 0.9%;
 PCT/US01/00663
 Score 12.8;
Pred. No. 1
 Mismatches
 .7e+02;
 Length 17;
 Indels
 0
 Gaps
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US-10-060-756A-1675/c
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 RESULT 248
 US-10-060-756A-146
 US-10-060-756A-144
 US-10-060-756A-146
 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 146
 Matches
 Query Match
 Publication No. US20 GENERAL INFORMATION:
 Sequence 1675, Application US/10060756A Publication No. US20030046717A1
 Query Match
Best Local S
 GENERAL INFORMATION
 Matches
 Sequence 146, A
Publication No.
 FILE REFERENCE: PB0177
CURRENT APPLICATION NUMBER: US/10/060,756A
CURRENT FILING DATE: 2002-01-30
 FILE REFERENCE: PB0177
CURRENT APPLICATION NUMBER: US/10/060,756A
CURRENT FILING DATE: 2002-01-30
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
 APPLICANT: Zhang, Jian
TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
 PRIOR FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/327,898
 PRIOR APPLICATION NUMBER: PCT/US01/00665 PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: PCT,
 TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
 PRIOR APPLICATION NUMBER: PCT/US01/00667
 LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
 NUMBER OF SEQ ID NOS: 480'
 PRIOR FILING DATE:
 Local
 Local Similarity
 APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00663
 APPLICATION NUMBER: US 09/864,761
 FILING DATE: 2001-01-30
 APPLICATION NUMBER: PCT/US01/00665
 1519 CACAAGCTGACCAAAC 1534
 1518 GCACAAGCTGACCAAA 1533
 14; Conservative
 14;
 N
 ING DATE:
 Similarity
 GCCCAAGCTCACCAAA 17
 CCCAAGCTCACCAAAC 16
 Application US/10060756A o. US20030046717A1
 Conservative
 2001-10-09
 001-01-30
 0.9%;
 0.9%;
87.5%;
PCT/US01/00668
 <u>.</u>
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 2;
 Mismatches
 <u>ب</u>
 Length 17;
 Length 17;
 Indels
 0,
 0,
 Gaps
 Gaps
 0
 0
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RESULT 249
US-10-060-756A-1677/c
; Sequence 1677, Application US/10060756A
; Sequence 1670, No. US20030046717A1
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 Query Match
Best Local Similarity
"hes 14; Conserve
 ; TYPE: DNA; Homo sapiens US-10-060-756A-1675
 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-060-756A-1677
 NUMBER OF SEQ ID NOS: 4804
SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 1677
 SEQ ID NO 1675
 Matches
 Query Match
Best Local
 CURRENT FILING DATE: 2002-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
 PRIOR FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/327,898
PRIOR FILING DATE: 2001-10-09
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 09/864,761
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
 APPLICANT: Zhang, Jian
TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
 SOFTWARE: Aeomica Sequence Listing Engine
 NUMBER OF SEQ ID NOS: 4804
 PRIOR FILING DATE:
PRIOR APPLICATION N
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
 CURRENT APPLICATION NUMBER: US/10/060,756A
CURRENT FILING DATE: 2002-01-30
 PRIOR FILING DATE: 2001-10-09
 PRIOR APPLICATION NUMBER: US 09/864,761
 PRIOR APPLICATION NUMBER: PCT/US01/00669
 FILE REFERENCE: PB0177
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00668
 LENGTH: 17
 ENGTH:
 Local
 2186 ATGTGATGAAAATAGC 2201
 2187 TGTGATGAAAATAGCA 2202
 16
 17 TGTTATAAAAATAGCA 2
 14;
 17
 Similarity
ATGTTATAAAAATAGC
 CATION NUMBER: US 60/327,898
 Conservative
 Conservative
 2001-05-23
 0.9%;
 0.9%;
 0;
 0;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8;
Pred. No. 1.
 Mismatches
 Mismatches
 1.7e+02;
 DB 1;
 DB 1;
 Length 17;
 Length 17;
 0,
 0,
 Gaps
 Gaps
 0,
 0,
```

US-10-060-998-588/c

Sequence 588, Application US/10060998 Publication No. US20030104530A1 GENERAL INFORMATION:

RESULT 250

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 RESULT 251
US-10-060-998-589/c
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 US-10-163-552-677
 ; TYPE: DNA; Homo sapiens US-10-060-998-588
 US-10-060-998-589
 CURRENT APPLICATION NUMBER: US/10/060,998
CURRENT FILING DATE: 2002-01-30
PRIOR APPLICATION UNMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 09/864,761
PRIOR FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/343,331
PRIOR APPLICATION NUMBER: US 60/343,331
PRIOR FILING DATE: 2001-12-21
PRIOR FILING DATE: 2001-12-21
 GENERAL INFORMATION:
 Sequence 589, Application US/10060998 Publication No. US20030104530A1 GENERAL INFORMATION:
 Sequence 677, Application US/10163552
Publication No. US20030105051A1
 SEQ ID NO 589
LENGTH: 17
 SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 588
 Matches
 Query Match
 Matches
 Query Match 0.9%;
Best Local Similarity 87.5%;
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic acid treatment
TITLE OF INVENTION: HER2
FILE REFERENCE: MBHB01-1.653-A (400/014)
CURRENT PPLICATION NUMBER: US/10/163,552
CURRENT FILING DATE: 2002-06-06
 CURRENT APPLICATION NUMBER: US/10/060,998
CURRENT FILING DATE: 2002-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR PPLICATION NUMBER: US 09/864,761
PRIOR PLING DATE: 2001-05-23
PRIOR PLING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/343,331
 SOFTWARE: Aeomica Sequence Listing Engine
 NUMBER OF SEQ ID NOS: 3056
 FILE REFERENCE: PB01108
 APPLICANT: Gu, Yizhong
TITLE OF INVENTION: HUMAN SODIUM-HYDROGEN EXCHANGER LIKE PROTEIN 1
 PRIOR FILING DATE: 2001-12-21
 NUMBER OF SEQ ID NOS: 3056
 FILE REFERENCE:
 TITLE OF INVENTION: HUMAN SODIUM-HYDROGEN EXCHANGER LIKE PROTEIN 1
 TYPE: DNA
ORGANISM: Homo sapiens
 ENGTH:
 Local
 1692 ATGGGAGTTTCCAAGA 1707
 1692 ATGGGAGTTTCCAAGA 1707
 16
 17
 14;
 14;
 Similarity
 ATGGCAGTTCCCAAGA 1
 ATGGCAGTTCCCAAGA 2
 Conservative
 Conservative
 0.9%;
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 0,
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
 Mismatches
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 of diseases or conditions related to leve
 DB 1;
 Length 17;
 Length 17;
 Indels
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 ; ORGANISM: Homo sapiens US-10-156-306-228
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 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-163-552-935
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 Query Match
Best Local Similarity
Matches 10; Conserve
 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-163-552-677
 US-10-156-306-228/c
 RESULT 253
 US-10-163-552-935/c
 SOFTWARE: PatentIn version 3.0
SEQ ID NO 228
LENGTH: 17
 Matches
 Sequence 228, Application US/10156306 Publication No. US20030119017A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 935
 Query Match
 Matches
 SEQ ID NO 677
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 Sequence 935, Application US/10163552 Publication No. US20030105051A1
 CURRENT APPLICATION NUMBER: US/10/156,306 CURRENT FILING DATE: 2002-05-28
 FILE REFERENCE: MBHB01-664-A (400/050)
 TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases TITLE OF INVENTION: Levels of IKK-Gamma and PKR
 FILE REFERENCE: MBHB01-1653-A (400/014)
CURRENT APPLICATION NUMBER: US/10/163,552
CURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 1997
 NUMBER OF SEQ ID NOS: 8013
 APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: McSwiggen, James
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic acid treatment
TITLE OF INVENTION: HER2
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, Jim
 SOFTWARE: PatentIn version 3.0
 TYPE: RNA
 LENGTH:
 ENGTH: 17
 2645 CTTCAGGAGATGATTC 2660
 2701 CCTCAGTATCCACACA 2716
 2317 CATCAGAGTGATGTCT 2332
 14;
 17 CCTCAGAATCCACAAA 2
 14;
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 Similarity
 CACCAGAGUGAUGUGU 17
 0.9%;
nilarity 87.5%;
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 Score 12.8; D. Pred. No. 1.7e 0; Mismatches
 Score 12.8;
Pred. No. 1
 Score 12.8;
Pred. No. 1
 Mismatches
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 1.7e+02;
 1.7e+02
 DB 1;
 of diseases or conditions related to level
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 or Conditions Relate
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US-10-156-306-1457/c
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 RESULT 256
US-10-156-306-365
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 S
 ; TYPE: RNA; Homo sapiens US-10-156-306-364
 US-10-156-306-364
 RESULT 257
 ; ORGANISM: Homo sapiens
US-10-156-306-365
 TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate TITLE OF INVENTION: Levels of IKK-Gamma and PKR

FILE REFERENCE: MBHB01-664-A (400/050)
CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
SOFTWARE: PatentIn version 3.0
SEQ ID NO 365
LENGTH: 17
 Sequence 1457, Application US/10156306 Publication No. US20030119017A1 GENERAL INFORMATION:
 Sequence 365, Applic Publication No. US20 GENERAL INFORMATION:
 Sequence 364, Application No. US200 GENERAL INFORMATION:
 Query Match 0.9%;
Best Local Similarity 62.5%;
Matches 10; Conservative
 Matches
 SEQ ID NO 364
 Query Match
Best Local
 TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate TITLE OF INVENTION: Levels of IKK-Gamma and PKR FILE REFERENCE: MBHB01-664-A (400/050) CURRENT APPLICATION NUMBER: US/10/156,306 CURRENT FILING DATE: 2002-05-28 NUMBER OF SEQ ID NOS: 8013
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases
TITLE OF INVENTION: Levels of IKK-Gamma and PKR
FILE REFERENCE: MBHB01-664-A (400/050)
CURRENT FILING DATE: 2002-05-28
CURRENT FILING DATE: 2002-05-28
CURBER OF SEQ ID NOS: 8013
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, James
 SOFTWARE: PatentIn version 3.0
 APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: McSwiggen, James
 TYPE: RNA
 ENGTH:
 Match 0.9%;
Local Similarity 62.5%;
les 10; Conservative
 2200 GCAGACTTTGGACTCG 2215
 2200 GCAGACTTTGGACTCG 2215
 N
 17
 1 ĠĠĀĠĀĊŪŪŪĠĠĀĊŪŪĠ 16
PatentIn version 3.0
 GGAGACUUUGGACUUG 17
 Application US/10156306 o. US20030119017A1
 Application US/10156306
 US20030119017A1
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
4; Mismatches 2;
 Score 12.8;
Pred. No. 1.
 Mismatches
 1.7e+02;
2;
 8; DB 1;
1.7e+02;
 Length 17;
 Length 17;
 0;
 or Conditions Relate
 0
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LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-156-306-1457
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 ; ORGANISM: Homo sapiens US-10-156-306-1545
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 ; ORGANISM: Homo sapiens US-10-156-306-2834
 RESULT 259
US-10-156-306-2834
 RESULT 258
US-10-156-306-1545
 Sequence 2834, Application US/10156306
publication No. US20030119017A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate.
TITLE OF INVENTION: Levels of IKK-Gamma and PKR
TITLE OF INVENTION: Levels of IKK-Gamma and PKR
TITLE REFERENCE: MHB01-664-A (400/050)
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 2834 LENGTH: 17
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 1545
 GENERAL INFORMATION:
 Matches 14;
 SEQ ID NO 1457
 Sequence 1545, Application US/10156306
Publication No. US20030119017A1
 Query Match 0.9%;
Best Local Similarity 87.5%;
 Matches
 Query Match
Best Local Similarity
 Matches
 Query Match
 TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment TITLE OF INVENTION: Levels of IKK-Gamma and PKR FILE REFERENCE: MEHBO1-664-A (400/050)
CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
 CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
 APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: McSwiggen, James
 TYPE: RNA
 TYPE: RNA
 LENGTH:
 LENGTH:
 Local Similarity
mes 9; Conserv
 2198 TAGCAGACTTTGGACT 2213
 2645 CTTCAGGAGATGATTC 2660
 16
 17
 10;
 1 UUGGAGACUUUGGACU 16
 ATAGCAGACTTTGGAC 2212
 CTTCAGATGATGATTC 1
 AUUGGAGACUUUGGAC 17
 Conservative
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 56.2%;
 0.9%;
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 <u>..</u>
 Score 12.8; DB 1;
Pred. No. 1.7e+02;
5; Mismatches 2
 Score 12.8;
Pred. No. 1
 Score 12.8;
Pred. No. 1.
 Mismatches
 Mismatches
 1.7e+02;
 .7e+02
 DB 1;
 DB 1;
 DB 1;
 Length 17;
 Length 17
 Length 17;
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 of Diseases
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 or Conditions Related
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 S
 US-09-263-959-616
 RESULT 262
 US-10-156-306-4950
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 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-156-306-4426
 ; ORGANISM: Homo sapiens
US-10-156-306-4950
 RESULT 261
 US-10-156-306-4426
 RESULT 260
 SEQ ID NO 4950
LENGTH: 17
TYPE: RNA
 Sequence 616, Application US/09263959
 Sequence 4950, Application US/10156306 Publication No. US20030119017A1 GENERAL INFORMATION:
 Patent No. US20020150891A1
GENERAL INFORMATION:
 Matches
 Query Match
Best Local Similarity
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 4426
 Sequence 4426, Application US/10156306 Publication No. US20030119017A1
 GENERAL INFORMATION:
APPLICANT: Hood, Leroy E.
APPLICANT: Rowen, Lee
APPLICANT: Roop, Ben F.
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI
NUMBER OF SEQUENCES: 1279
 Query Match
 Matches
 GENERAL INFORMATION:
 TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate TITLE OF INVENTION: Levels of IKK-Gamma and PKR FILE REFERENCE: MBHB01-664-A (400/050)
CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
 CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
SOFTWARE: Patentin version 3.0
 APPLICANT: McSwiggen, James
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of IKK-Gamma and PKR
FILE REFERENCE: MBHB01-664-A (400/05)
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
 APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, James
 ENGTH: 17
STREET:
 Local Similarity 75.0
 ADDRESSEE:
 1574 CCAGCTCCTCCATGAA 1589
 1574 CCAGCTCCTCCATGAA 1589
 12; Conservative
 1 CCAGCUCUUCCAAGAA 16
 CCAGCUCUUCCAAGAA 17
6300 Columbia Center,
 Seed and Berry LLP
 0.9%;
 0.9%; Score 12.8; I
75.0%; Pred. No. 1.70
ntive 2; Mismatcheв
 2;
 Score 12.8;
Pred. No. 1
 Mismatches
 701 Fifth Avenue
 .7e+02;
 DB 1; Length 17; .7e+02;
 DB 1;
 Length 17;
 Indels
 0
 0;
 0
 0
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밁
 RESULT 263
US-09-263-959-638
 Ś
 US-09-263-959-616
 0.9%;
Best Local Similarity 92.9%;
Matches 13; Conservative
 Sequence 638, Application US/09263959 Patent No. US20020150891A1
 TELEFAX: (206) 682-6031 INFORMATION FOR SEQ ID NO:
 GENERAL INFORMATION:
 TELEFAX: (206) 682-60 INFORMATION FOR SEQ ID NO:
 COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE: 05-MAR-1999
CLASSIFICATION:
 ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 APPLICANT:
APPLICANT:
 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
 SEQUENCE CHARACTERISTICS:
LENGTH: 14 base pairs
 TELECOMMUNICATION INFORMATION: TELEPHONE: (206) 622-4900
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
 SEQUENCE CHARACTERISTICS:
 ATTORNEY/AGENT INFORMATION:
 CORRESPONDENCE ADDRESS:
 APPLICANT: Rowen, Lee
APPLICANT: KOOP, BEN F.
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI
NUMBER OF SEQUENCES: 1279
 TELECOMMUNICATION INFORMATION:
 ATTORNEY/AGENT INFORMATION:
 NAME: McMasters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920010.426C2
 NAME: McMasters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920010.426C2
 STREET: 6300 (CITY: Seattle
 TELEPHONE:
 COUNTRY:
 STATE:
 STRANDEDNESS:
 TYPE: nucleic acid
 CLASSIFICATION:
 APPLICATION NUMBER: FILING DATE: 05-MAR
 COUNTRY:
STRANDEDNESS:
 ADDRESSEE:
 TELEPHONE:
 ropology:
 LENGTH: 14 base pairs
 1882 ATGATGAAGATGAT 1895
 98104-7092
 Washington
 6300 Columbia Center, 701 Fifth Avenue
 S
 (206) 682-6031
 Hood, Leroy E.
 linear
 Seed and Berry LLP
 (206) 622-4900
 05-MAR-1999
 US/09/263,959
 616:
 0
 Score 12.4; DB 1;
Pred. No. 1.3e+02;
 Mismatches
 Length 14;
 Indels
 0
 Gaps
 0
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US-09-263-959-638

TOPOLOGY:

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 US-10-077-508-5/c
 US-10-077-508-3
 US-10-077-508-3
 GENERAL INFORMATION:
 SEQ ID NO 5
 Sequence 5, Application US/10077508 Publication No. US20030138784A1
 Matches
 Query Match
Best Local Similarity
 SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 3
 Sequence 3, Application US/10077508 Publication No. US20030138784A1
 Best Local Similarity Matches 13; Conserv
 Publication No. US20 GENERAL INFORMATION:
 Query Match
 CURRENT FILING DATE: 2002-02-15
PRIOR APPLICATION NUMBER: 09/53,366
PRIOR FILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: 60/125,864
PRIOR FILING DATE: 1999-03-24
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver: 2.1
 FILE REFERENCE: UTSD:557
CURRENT APPLICATION NUMBER: US/10/077,508
CURRENT FILING DATE: 2002-02-15
 APPLICANT: JOHNSTON, STEPHEN A.
TITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS
 APPLICANT: SYKES, APPLICANT: JOHNS
 PRIOR APPLICATION NUMBER: 60/125,864
PRIOR FILING DATE: 1999-03-24
NUMBER OF SEQ ID NOS: 5
 CURRENT APPLICATION NUMBER: US/10/077,508 CURRENT FILING DATE: 2002-02-15
 APPLICANT: JOHNSTON, STEPHEN A.
TITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS
FILE REFERENCE: UTSD:557
 APPLICANT: SYKES, APPLICANT: JOHNS
 PRIOR FILING DATE: 2000-03-24
 PRIOR APPLICATION NUMBER: 09/535,366
 FEATURE: OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Primer
 LENGTH: 14
TYPE: RNA
ORGANISM: Artificial Sequence
OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Primer
 ORGANISM: Artificial Sequence FEATURE:
 ENGTH:
 1882 ATGATGAAGATGAT 1895
 1882 ATGATGAAGATGAT 1895
 14
 9
 AUGAUGAUGAUGAU 14
 Conservative
 Conservative
 KATHRYN F.
 KATHRYN F.
 0.9%;
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 Score 12.4; DB 1;
Pred. No. 1.3e+02;
 Score 12.4; DB 1;
Pred. No. 1.3e+02;
 Mismatches
 Length 14
 Length 14;
 Indels
 Synthetic
 Synthetic
 0
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 Gaps
 Gaps
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 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-277-494-74
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 US-10-077-621-3
 US-10-077-621-3
 RESULT 267
 US-10-277-494-74
 RESULT 266
 Query Match
Best Local S
Matches 13
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 74 LENGTH: 14
 Sequence 3, Application US/10077621 Publication No. US20020146733A1
 Matches
 SEQ ID NO 3
 GENERAL INFORMATION
 Query Match
Best Local (
 GENERAL
 Sequence 74, Application US/10277494 Publication No. US20030186909A1
 Best Local Similarity
Matches 9; Conserv
 Query Match
 Matches
 CURRENT APPLICATION NUMBER: US/10/077,621
CURRENT FILING DATE: 2002-02-15
PRIOR APPLICATION NUMBER: 09/535,366
PRIOR FILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: 60/125,864
PRIOR FILING DATE: 1999-03-24
NUMBER OF SEQ ID NOS: 5
 CURRENT APPLICATION NUMBER: US/10/277,494
CURRENT FILING DATE: 2002-10-21
NUMBER OF SEQ ID NOS: 446
 APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or conditions Related To Level
TITLE OF INVENTION: Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-K (400/064)
 SOFTWARE: PatentIn Ver. 2.1
 APPLICANT: JOHNSTON, STEPHEN A.
TITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS
FILE REFERENCE: UTSD:557
 APPLICANT: SYKES, APPLICANT: JOHNS
 FEATURE: OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Primer
 TYPE: RNA
ORGANISM: Artificial Sequence
 LENGTH: 14
 y Match 0.9%;
Local Similarity 71.4%;
hes 10; Conservative
1882 ATGATGAAGATGAT 1895
 2317 CATCAGAGTGATGT 2330
 1882 ATGATGAAGATGAT 1895
 INFORMATION:
 14
 0.9%;
il Similarity 92.9%;
13; Conservative
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 Conservative
 KATHRYN F
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 0
 Score 12.4; DB 1;
Pred. No. 1.3e+02;
 Score 12.4;
Pred. No. 1.
 Score 12.4;
Pred. No. 1
 Mismatches
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 Mismatches
 1.3e+02;
 .3e+02;
 DB 1;
 DB 1;
 Length 14;
 Length 14;
 Length 14;
 Indels
 Indels
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 Synthetic
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 Gaps
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RESULT 270
US-10-077-232-5/c
, Sequence 5, Application US/10077232
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 Query Match
Best Local Similarity
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 ; OTHER INFORMATION: Description of Artificial Sequence: ; OTHER INFORMATION: Primer US-10-077-232-3
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 S
 US-10-077-621-5
 US-10-077-621-5/c
 Sequence 3, Application US/10077232
Publication No. US20020150940A1
GENERAL INFORMATION:
APPLICANT: SYKES, KATHRYN F.
APPLICANT: JOHNSTON, STEPHEN A.
 SEQ ID NO 3
 Sequence 5, Application US/10077621 Publication No. US20020146733A1
 Matches 13; Conservative
 Query Match
 GENERAL INFORMATION:
 PRIOR APPLICATION NUMBER: 60/125,864
PRIOR FILING DATE: 1999-03-24
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver. 2.1
 FILE REFERENCE: UTSD:557
CURRENT APPLICATION NUMBER: US/10/077,232
CURRENT FILING DATE: 2002-02-15
 CURRENT APPLICATION NUMBER: US/10/077,621
CURRENT FILLING DATE: 2002-02-15
PRIOR APPLICATION NUMBER: 09/535,366
PRIOR FILING DATE: 2000-03-24
 PRIOR APPLICATION NUMBER: 09/535,366 PRIOR FILING DATE: 2000-03-24
 APPLICANT: JOHNSTON, STEPHEN A.
TITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS
 APPLICANT: JOHNSTON, STEPHEN A.
TITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS
FILE REFERENCE: UTSD:557
 NUMBER OF SEQ ID NOS: 5
SOFTWARE: PatentIn Ver. 2.1
 PRIOR APPLICATION NUMBER: 60/125,864
PRIOR FILING DATE: 1999-03-24
 APPLICANT: SYKES, APPLICANT: JOHNST
 ORGANISM: Artificial Sequence
 LENGTH: 14
TYPE: RNA
 FEATURE:
 OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Primer
 TYPE: RNA
ORGANISM: Artificial Sequence
 FEATURE:
 Q ID NO 5
LENGTH: 14
 Match 0.9%;
Local Similarity 92.9%;
 1882 ATGATGAAGATGAT 1895
 1882 ATGATGAAGATGAT 1895
 14
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 Conservative
 KATHRYN F
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 Pred. No. 1.3e
4; Mismatches
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 Score 12.4;
Pred. No. 1
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 Length 14;
 Indels
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 Length 14;
 Synthetic
 Synthetic
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 Query Match
Best Local Similarity
"Artches 9; Conserva
 ; OTHER INFORMATION: Description of Artificial Sequence: ; OTHER INFORMATION: Primer US-10-077-247-3
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 US-10-077-232-5
 US-10-077-247-3
 PRIOR FILING DATE: 1999-03-24
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 3
 SEQ ID NO 5
LENGTH: 14
TYPE: RNA
ORGANISM: Artificial Sequence
 Sequence 3, Application US/10077247 Publication No. US20020155508A1
 GENERAL INFORMATION:
 Matches
 Best Local Similarity
 Query Match
 Publication No. US20020150940A1 GENERAL INFORMATION:
 TITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS FILE REFERENCE: UTSD:557
CURRENT APPLICATION NUMBER: US/10/077,247
CURRENT FILING DATE: 2002-02-15
PRIOR APPLICATION NUMBER: 09/535,366
PRIOR PILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: 60/125,864
 CURRENT APPLICATION NUMBER: US/10/077,232
CURRENT FILING DATE: 2002-02-15
PRIOR APPLICATION NUMBER: 09/535,366
PRIOR FILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: 60/125,864
PRIOR FILING DATE: 1999-03-24
NUMBER OF SEQ ID NOS: 5
 APPLICANT: JOHNSTON, STEPHEN A.
TITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS
FILE REFERENCE: UTSD:557
 APPLICANT: SYKES, KATHRYN F.
APPLICANT: JOHNSTON, STEPHEN A.
 APPLICANT: SYKES, KATHRYN F. APPLICANT: JOHNSTON, STEPHE
 SOFTWARE: PatentIn Ver. 2.1
 ORGANISM: Artificial Sequence
 TYPE: RNA
 OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Primer
 FEATURE:
 ENGTH: 14
 FEATURE:
 1882 ATGATGAAGATGAT 1895
 1882 ATGATGAAGATGAT 1895
 14 ATGATGATGAT 1
 13;
AUGAUGAUGAUGAU 14
 Conservative
 Conservative
 0.9%;
92.9%;
 0.9%;
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 Score 12.4;
Pred. No. 1
 Score 12.4; DB 1;
Pred. No. 1.3e+02;
 Mismatches
 Mismatches
 1.3e+02
 .3e+02
 DB 1;
 Length 14;
 Synthetic
 Synthetic
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 Gaps
 Gaps
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RESULT 272 US-10-077-247-5/c

Sequence 5, Application US/10077247 Publication No. US20020155508A1

KATHRYN F.

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US-10-077-392-5/c
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 FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: Primer
US-10-077-247-5
 RESULT 274
 US-10-077-392-3
 US-10-077-392-3
 RESULT 273
 Sequence 5, Application US/10077392 Publication No. US20020160402A1 GENERAL INFORMATION:
 Sequence 3, Application US/10077392 Publication No. US20020160402A1
 Matches
 Query Match
Best Local
 SEQ ID NO 3
 GENERAL INFORMATION:
 Matches
 SEQ ID NO 5
 Query Match
 PRIOR APPLICATION NUMBER: 09/535,366
PRIOR FILING DATE: 2000-03-24
PRIOR PPLICATION NUMBER: 60/125,864
PRIOR FILING DATE: 1999-03-24
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver. 2.1
 CURRENT FILING DATE: 2002-02-15
PRIOR APPLICATION NUMBER: 09/535,366
PRIOR FILING DATE: 2000-03-24
 FILE REFERENCE: UTSD:557
CURRENT APPLICATION NUMBER: US/10/077,392
CURRENT FILING DATE: 2002-02-15
 PRIOR APPLICATION NUMBER: 60/125,864
PRIOR FILING DATE: 1999-03-24
NUMBER OF SEQ ID NOS: 5
 APPLICANT: SYKES, APPLICANT: JOHNST
 APPLICANT: JÖHNSTON, STEPHEN A.
TITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS
 APPLICANT: SYKES, APPLICANT: JOHNS
 SOFTWARE: PatentIn Ver. 2.1
 FILE REFERENCE: UTSD:557
CURRENT APPLICATION NUMBER: US/10/077,247
 APPLICANT: JOHNSTON, STEPHEN A.
TITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS
FILE REFERENCE: UTSD: 557
 LENGTH: 14
TYPE: RNA
ORGANISM: Artificial Sequence
 OTHER INFORMATION:
 FEATURE:
 ORGANISM: Artificial Sequence
 TYPE: RNA
 PPLICANT: JOHNSTON, STEPHEN A.
ITLE OF INVENTION: LINEAR AND CIRCULAR EXPRESSION ELEMENTS
 ENGTH: 14
 Local
 Local
 1882 ATGATGAAGATGAT 1895
 1882 ATGATGAAGATGAT 1895
 14 ATGATGATGAT 1
 13;
 H
 0.9%;
Similarity 92.9%;
 Similarity 9; Conserv
 AUGAUGAUGAUGAU 14
 Conservative
 Conservative
 KATHRYN F.
 KATHRYN F.
 Description of Artificial Sequence: Synthetic Primer
 0.9%;
 0,
 Score 12.4; DB 1;
Pred. No. 1.3e+02;
4; Mismatches 1;
 Score 12.4; DB 1;
Pred. No. 1.3e+02;
 Mismatches
 1.3e+02;
 Length 14;
 Length 14;
 Indels
 Indels
 Synthetic
 0
 <u>,,</u>
 Gaps
 Gaps
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 ; OTHER INFORMATION: Description of Artificial Sequence: Probe US-09-771-933-174
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 US-10-077-392-5
 RESULT 276
US-09-848-754A-9628
 RESULT 275
US-09-771-933-174/c
 Sequence 9628, Application US/09848754A

Publication No. US20030073207A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FLING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
SOFTWARE: PatentIn version 3.0
 Query Match
Best Local Similarity
Matches 13; Conserv
 SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 174
 GENERAL INFORMATION:
 SEQ ID NO 5
 SEQ ID NO 9628
 Matches
 Publication No.
 Sequence 174,
 Query Match
 CURRENT APPLICATION NUMBER: US/09/771,933
CURRENT FILING DATE: 2001-01-30
NUMBER OF SEQ ID NOS: 205
 APPLICANT: Gill-Garrison, Rosalynn D
APPLICANT: Martin, Christopher J
APPLICANT: Sanchez-Felix, Manuel V
TITLE OF INVENTION: Computer-assisted Means
TITLE OF INVENTION: Factors
 SOFTWARE: PatentIn Ver. 2.1
 PRIOR FILING DATE: 199
 CURRENT APPLICATION NUMBER: US/10/077,392
CURRENT FILING DATE: 2002-02-15
 FILE REFERENCE: 620-130
 PRIOR APPLICATION NUMBER: 60/125,864
 PRIOR APPLICATION NUMBER: 09/535,366
PRIOR FILING DATE: 2000-03-24
 ORGANISM: Artificial Sequence FEATURE:
 TYPE: DNA
 OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Primer
 ORGANISM: Artificial Sequence
 FEATURE:
 TYPE: RNA
 LENGTH:
LENGTH: 15
 ENGTH:
 Local Similarity
 1880 AGATGATGAAGATG 1893
 1882 ATGATGAAGATGAT 1895
 14
 14
 5
 AGATGATGGAGATG
 Application US/09771933
 Conservative
 Conservative
 US20030023387A1
 1999-03-24
 92.9%;
 0.9%;
 <u>.</u>
 0
 Score 12.4;
Pred. No. 1.
 Score 12.4;
Pred. No. 1.
 Mismatches
 Mismatches
 ; DB 1;
L.3e+02;
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 for
 DB 1;
 Assessing
 Length 15;
 Length 14;
 Indels
 Synthetic
 Lifestyle Risk
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 ; SEQUENCE DESCRIPTION: SEQ ID NO: 157: US-09-943-983-157
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 RESULT 277
US-09-943-983-157/c
 ; OTHER INFORMATION: Description of Artificial Sequence: Enzymatic Nucleic acid US-09-848-754A-9628
 Matches
 Query Match
Best Local Similarity
 Sequence 157, Applicat Publication No. US2003 GENERAL INFORMATION:
 Query Match
Best Local Similarity
 Matches
 NAME: KAMMERER, PATRICIA A.
REGISTRATION NUMBER: 29,775
REFERENCE/DOCKET NUMBER: INNS:008
INFORMATION FOR SEQ ID NO: 157:
 TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
 APPLICATION NUMBER: 08/913,833
FILING DATE: 1997-09-15
APPLICATION NUMBER: EP 96870005.4
FILING DATE: 26 Jan 1996
APPLICATION NUMBER: EP 96870081.5
FILING DATE: 25 Jun 1996
ATTORNEY/AGENT INFORMATION:
 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 6.0 / F
CURRENT APPLICATION DATA:
 1856 TTTCTGATCTGGTG 1869
 TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
 SEQUENCE CHARACTERISTICS:
 NUMBER OF SEQUENCES: 164
CORRESPONDENCE ADDRESS:
ADDRESSE: ARNOLD, WHITE & DURKEE STREET: P.O. BOX 4433
 2317 CATCAGAGTGATGT 2330
 ANTI-SENSE: NO
 HYPOTHETICAL: NO
 PRIOR
 ROSSAU, RUDI
TITLE OF INVENTION: METHOD FOR DETECTION OF DRUG-INDUCED
 APPLICANT: STUYVER, LIEVEN LOUWAGIE, JOOST
 10; Conservative
 2 CACCAGAGUGAUGU 15
 TYPE: nucleic acid
STRANDEDNESS: single
 APPLICATION NUMBER: US/09/943,983 FILING DATE: 31-Aug-2001 APPLICATION DATA:
 CITY: HOUSTON
STATE: TEXAS
 COUNTRY: USA
ZIP: 77210-4433
 LENGTH: 15 base pairs
 Application US/09943983
o. US20030077575A1
 Conservative
 92.9%;
 0.9%;
 MUTATIONS IN THE REVERSE TRANSCRIPTASE GENE
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 Score 12.4; DB 1;
Pred. No. 1.5e+02;
0; Mismatches 1;
 Score 12.4; DB 1;
Pred. No. 1.5e+02;
 Mismatches
 ASCII text output
 DB 1;
 Length 15;
 Length 15;
 Indels
 Indels
 0,
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 Gaps
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 ; TYPE: RNA; Homo sapiens US-10-440-850-911
 RESULT 279
US-10-440-850-911/c
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 ; LENGTH: 15
; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-277-494-57
 RESULT 278
US-10-277-494-57
; Sequence 57, Application US/10277494
; Publication No. US20030186909A1
 Query Match
Best Local Similarity
Matches 13; Conserva
 SEQ ID NO 911
 Sequence 911, Application US/10440850 Publication No. US20030207837A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 57
 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
 Matches
 Query Match
 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Induction of Graft Tolerance
TITLE OF INVENTION: Method and Reagent for the Induction of Graft Tolerance
TITLE OF INVENTION: Method and Reagent for the Induction of Graft Tolerance
TITLE OF INVENTION: Method and Reagent for the Induction of Graft Tolerance
FILE REFERENCE: 250/130 (MSHB00-900-A)
CURRENT FILING DATE: 2003-05-19
PRIOR APPLICATION NUMBER: US/09/650,012
PRIOR FILING DATE: 1906-01-12
PRIOR FILING DATE: 1996-01-12
PRIOR FILING DATE: 1996-01-07
PRIOR FILING DATE: 1995-07-07
PRIOR FILING DATE: 1995-07-07
PRIOR FILING DATE: 1995-07-07
 CURRENT APPLICATION NUMBER: US/10/277,494
CURRENT FILING DATE: 2002-10-21
NUMBER OF SEQ ID NOS: 446
 TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or conditions Related To Level TITLE OF INVENTION: Epidermal Growth Factor Receptors FILE REFERENCE: MBHB00-958-K (400/064)
 PRIOR FILING DATE: 1998-03-11
NUMBER OF SEQ ID NOS: 2285
SOFTWARE: PatentIn version 3.0
 APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: Stinchcomb, Dan
 APPLICANT:
 PRIOR APPLICATION NUMBER: US 09/038,073
 LENGTH: 15
 Match 0.9%;
Local Similarity 71.4%;
 1887 GAAGATGATTGGGA 1900
 2317 CATCAGAGTGATGT 2330
14 GAAGATGATGGGGA 1
 10;
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 CACCAGAGUGAUGU 15
 Jarvis, Thale
 Conservative
 Conservative
 0.9%; Score 12.4;
92.9%; Pred. No. 1.
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 Score 12.4;
Pred. No. 1.
 Mismatches
 Mismatches
 1.5e+02;
 1.5e+02
 Length 15;
 Indels
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RESULT 280
US-10-097-175-50
US-10-097-175-50
; Sequence 50, Application US/10097175
; Publication No. US20030045680A1

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US-10-287-919-2272/c
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 ; LENGTH: 15
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-010-802-192
 US-10-010-802-192
 RESULT 281
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 RESULT 282
 ; OTHER INFORMATION: Androgen Responsive Element US-10-097-175-50
 Sequence 2272, Application US/10287919 Publication No. US20030085830A1
 SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 192
 GENERAL INFORMATION:
 Query Match 0.9%;
Best Local Similarity 92.9%;
Matches 13; Conservative
 Matches
 Query Match
Best Local :
 SEQ ID NO 50
 Publication No.
 APPLICANT: Stephens, J. Claiborne
APPLICANT: Windemuth, Andreas
TITLE OF INVENTION: Drug Target Isogenes: Polymorphisms in the Interleukin
TITLE OF INVENTION: 4 Receptor Alpha Gene
FILE REFERENCE: MMH-0002US2 IL4R alpha
CURRENT APPLICATION UNMEER: US/10/010,802
CURRENT FILING DATE: 2001-11-09
PRIOR APPLICATION NUMBER: US/10/010,802
PRIOR APPLICATION NUMBER: US/10/010,802
PRIOR APPLICATION NUMBER: US/10/010,802
PRIOR APPLICATION NUMBER: US/10/010,802
PRIOR APPLICATION NUMBER: US/10/010,904
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 413
 APPLICANT:
 PRIOR APPLICATION NUMBER: 60/352,399
PRIOR FILING DATE: 2002-01-28
NUMBER OF SEQ ID NOS: 102
SOFTWARE: FastSEQ for Windows Version 4.0
 APPLICANT: FINDEIS, MARK A.

TITLE OF INVENTION: PEPTIDIC MODULATORS OF THE ANDROGEN RECEPTOR FILE REFERENCE: PPI-110

CURRENT APPLICATION NUMBER: US/10/097,175

CURRENT FILING DATE: 2002-03-12

PRIOR APPLICATION NUMBER: 60/275,240

PRIOR FILING DATE: 2001-03-12

PRIOR FILING DATE: 2001-03-12
APPLICANT: Feldmann,
 APPLICANT:
 APPLICANT: Genaissance Pharmaceuticals
 LENGTH: 15
TYPE: DNA
ORGANISM: Artificial Sequence
 APPLICANT:
 Local
 1543 CTGCGGAGACAGGT 1556
 2633 GAAGTTCTTGTTCT 2646
 13;
 _
 2 GAAGTTCCTGTTCT 15
 Similarity
 CTGTGGAGACAGGT 14
 Nandabalan, Krishnan
 Denton,
 Chew, Anne
 MUELLER, JOHN
 Application US/10010802
b. US20030078220A1
 Conservative
 VIBHA B.
 JOHN L.
 R. Rex
Richard J.; Global Determinants,
 92.9%;
 Score 12.4; Di
Pred. No. 1.5e
0; Mismatches
 0
 Score 12.4;
Pred. No. 1
 Mismatches
 .5e+02;
 DB 1;
 5e+02
 DB 1;
 Length 15;
 Length 15
 Indels
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 Gaps
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 ; I.OCATION: (1403144)...(1403158); OTHER INFORMATION: Chromosome = 1 US-10-287-919-2272
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 RESULT 284
US-09-837-992-29
 US-10-287-919-2400
 RESULT 283
US-10-287-919-2400/c
 GENERAL INFORMATION:
APPLICANT: Feldmann, Richard J.; Global Determinants,
APPLICANT: Feldmann, Richard J.; Global Determinants,
TITLE OF INVENTION: Methanococcus jannaschii complete
FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333
CURRENT APPLICATION NUMBER: US/10/287,919
CURRENT FILING DATE: 2002-11-05
NUMBER OF SEQ ID NOS: 2706
SOFTWARE: Proprietary
 NUMBER OF SEQ ID NOS: 2706
SOFTWARE: Proprietary
SEQ ID NO 2272
LENGTH: 15
 SEQ ID NO 2400
LENGTH: 15
 Matches
 Sequence 2400, Application US/10287919 Publication No. US20030085830A1
 Matches
 APPLICANT: Tian,
APPLICANT: Schul
APPLICANT: Shan,
APPLICANT: Tular
 GENERAL INFORMATION:
 Sequence 29, Application US/09837992 Patent No. US20020081687A1
 Best Local Similarity Matches 13; Conserv
 Query Match
 Query Match
Best Local Similarity
 TITLE OF INVENTION: Sitssterolemia Susceptibility Gene (SSG):
TITLE OF INVENTION: and Methods of Use
FILE REFERENCE: 018781-006020US
CURRENT APPLICATION NUMBER: US/09/837,992
CURRENT FILING DATE: 2001-04-18
PRIOR APPLICATION NUMBER: US 60/198,465
PRIOR PELLING DATE: 2000-04-18
PRIOR PILING DATE: 2000-04-18
PRIOR APPLICATION NUMBER: US 60/204,234
PRIOR FILING DATE: 2000-05-15
 TITLE OF INVENTION: Methanococcus jannaschii complete genome. FILE REFERENCE: Jim Zegeer Law Offices - 703-684-8333 CURRENT APPLICATION NUMBER: US/10/287,919 CURRENT FILING DATE: 2002-11-05
NUMBER OF SEQ ID NOS: 45
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 Tularik Inc.
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Pred. No. 1.
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Pred. No. 1.5e+02;
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Sequence 26, Application US/09816460C
Publication No. US20030087235A1
GENERAL INFORMATION:
APPLICANT: Dairkee, Shanaz H.
APPLICANT: Li, Zheng
TITLE OF INVENTION: PROGNOSTIC METHODS FOR BREAST CANCER
FILE REFERENCE: CPMC-010/00US
CURRENT APPLICATION NUMBER: US/09/816,460C
CURRENT APPLICATION NUMBER: US/09/816,460C
CURRENT FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 47
SOFTWARE: Patentin version 3.1
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COTHER INFORMATION: synthetic primer
US-09-816-460C-26
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; OTHER INFORMATION: 5' splicing site for exon
US-09-837-992-29
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Run

9

OM nucleic -

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 2 summaries
 Total number of hits satisfying chosen
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 12 bp mRNA linear EST 06-DEC E012340w-024-010-G19-SP6 MPIZ-ADIS-024-leaf Beta vulgaris cDNA clone 024-010-G19 5-PRIME, mRNA secuence
 Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes Plant J. 32 (5), 845-857 (2002)

Contact: Weisshaar B
 Beta vulgaris
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Caryophyllales; Amaranthaceae; Beta.
 Carl-von-Linne Weg 10, 50829 Koeln, Fax: 00492215062851
 ADIS DNA core facility at MPIZ Max-Planck-Institute for Plant Breeding Research
 BQ587288
BQ587288.1 GI:26116870
 Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H
 Beta vulgaris
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 (bases 1 to 12)
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 IDENTITY NUC
 us-09-954-556-3
 December
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 Copyright
weisshaa@mpiz-koeln.mpg.de
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12
 GenCore version 5.1.6 (c) 1993 - 2003 Compugen Ltd.
 2003, 12:04:41; Search time 0.001 Seconds (without alignments) 67.392 Million cell updates/

 BQ587288
BQ587706
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 cell updates/sec
 ACCESSION: BQ587788
ACCESSION: BQ587706
 Description
 EST 06-DEC-2002
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AUTHORS
 RESULT 2
BQ587706/c
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 Ş
 FEATURES
 COMMENT
 ORGANISM
 ACCESSION
 DEFINITION
 BASE COUNT
 FEATURES
 KEYWORDS
 VERSION
 TITLE
 Matches
 Query Match
 Best Local Similarity
 JOURNAL
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 12
 11;
 Beta vulgaris
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Result

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Score 10.4 10.4

Database

Minimum

DB DB

eq geq

Scoring table: Sequence: Perfect score: Title:

VERSION KEYWORDS

ORGANISM

ACCESSION DEFINITION

REFERENCE

AUTHORS

COMMENT

JOURNAL

TITLE

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 ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
 Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes Plant J. 32 (5), 845-857 (2002) Contact: Weisshaar B
 Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehand Radelof,U.
 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Caryophyllales; Amaranthaceae; Beta.
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В 8